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Title:
Perfect score:
Sequence:
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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seq length: 2000000000
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Compugen Ltd.
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Pred. No. 18 the number of results predicted by chance to have

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No. იიი იიიი 0000 0000 278 214.4 214.4 214.4 214.4 210.2 210.2 200.5 20 518.4 493.4 493.4 466.8 465.8 411.4 405.4 135 119.4 119.4 119.4 119.4 73.4 73.4 73.4 73.4 73.2 70.2 325.4 321.8 321.8 321.8 316.4 Query 45256 137334 171595 171595 209366 213593 273571 321 207 207 207 207 207 207 207 207 6283 6283 6283 6283 6283 6283 6283 59554 97075 151620 188460 269 Length DB AC018589 AC010542 BD030595 BD139357 AX330610 AX330791 AX331008 AX408097 AX252053 AC018557 AC018551 AC018551 AC018551 AX348836 AX252053 AX34430 AC1212851 AC12131402 AC131402 AX061665 AF135381 G30204 AX695587 AC010289 BD085905 AR275027 BD085952 BD023892 AF135380 AF145216 BC004380 AF096895 AR306550 AX061624 AR306591 AF057306 BD085905 Elongatio AR275027 Sequence BD085952 Elongatio BD023892 Sequence AP135380 Homo sapi AF145216 Homo sapi BC004380 Homo sapi BC004380 Homo sapi BC004381 Homo sapi AR061665 Sequence AF135381 Homo sapi AR30616591 Sequence AF135381 Homo sapi G30204 human STS Sexuence AF010289 Homo sapi G30204 Homo sapi G30204 Homo sapi BD030595 Sequence BD139357 Extended AX330610 Sequence AX330610 Sequence AX331008 Sequence AX331008 Sequence AX331008 Sequence AX079435 Sequence AC01857 Homo sapi AF253064 Rattus no AX252053 Sequence AX34431 Sequence AX34836 Sequence AY047360 Mus muscu AF253065 Rattus no AX252052 Sequence AX344430 Sequence AX344430 Sequence AX344430 Sequence AX346835 Sequence AX346835 Sequence AC0212952 Mus muscu AC121275 Mus muscu AC1212951 Mattus no AC131402 Rattus no AC137048 Rattus no AF401531 Mus muscu AC137048 Rattus no AF401531 Mus muscu AC137048 Rattus no AR306550 Sequence AX061624 Sequence AF096895 Homo Description

	Xia, D., Rui, M., Tang, J. and Ma, D.
11,Q., Sun,R.,	Gui, L., Tang, Y., Li, F., Song, Q., Di, C., Wang, L., Sl
Gu, W., Huang, J	Han, W., Lou, Y., Tang, J., Zhang, Y., Chen, Y., Li, Y.,
	1 (bases 1 to 530)
:; Homo.	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Euteleostomi;	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
	Homo sapiens
	Homo sapiens (human)
	FLI_CDNA.
	AF096895.2 GI:9989691
	AF096895
complete cds.	Homo sapiens chemokine-like factor 1 (CKLF1) mRNA, complete cds.
PRI 18-JUL-20	AF096895 530 bp mRNA linear

PRI 18-JUL-2001

Huang, J.,

ALIGNMENTS

RESULT 1
AF096895
LOCUS
DEFINITION
ACCESSION

VERSION KEYWORDS

SOURCE ORGANISM

REFERENCE AUTHORS

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REMARK
COMMENT
FEATURES
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TITLE
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Sequence update by submitter
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Biochem. J. 357 (Pt 1),
21308461
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Submitted (07-SEP-2000)
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Han, W.L., Li, Y., Zhang, Y.M.,
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                          TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTTGTGCATGAAA 434
                                                                        TTGGTGGAGGGGTTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                          CAGTATTCATGCTCATCGTATCTGTGTTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                                          TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
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/product="chemokine-like factor 1"
/product="chemokine-like factor 1"
/protein_id="AAF06722.1"
/db_xref="GI:6288734"
/translation="MDNVQPKH, KHRPFCFSVKGHVKMLRLDIINSLVTTVFMLIVSVL
ALIPETTTLTVGGGVFALVTANCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
108 c 137 g 129 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression inhibited
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"

/mol type="mRNA"

/db xref="taxon:9606"

/cell_line="U937"

/note="exposed to phytohaemagglutinin (PHA)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note="increased expression in expression inhibited by IL-10"
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99.8%;
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RESULT 3
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Edwards, J.-B.D.M., Bougueleret, L. and
Edwards, J.-B.D.M.'s encoding proteins
Complementary DNA's encoding proteins
Patent: US 6548633-A 29 15-APR-2003;
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                                                                                                     AAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                                                                                                                       TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAAGCCTGTGCATGAAA
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                                   TCTGTATTCTTCCAAAAAAAAAAAAAAAA 522
                                                                      AAAAAGAAGTTTTGTAATTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATT
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polyA_site
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AX061624.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                             AAAAAGAAGTTTTGTAATTTTATATTTACTTTTTTAGTTTTGATACTAAGTATTTAAACATATT
                                                                                                                                             TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGGCCTGTGCATGAAA
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                                                             AAAAAGAAGTTTTGTAATTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATT
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/db_xref="G1:12406710"
/taanslation="MONVOPKIKHRPECFSVKGHVKMLRLDIINSLVTTVFMLIVSVL
ALIPETTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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/mol_type="genomic_DNA"
/db_xref="taxon:9606"
144. .443
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99.4%;
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PR 13.NOV-1998 UP 2000521191
PR 13.NOV-1997 US 60/06677,17-DEC-1997 US 60/06957 PR
10-AUG-1998 US 60/06677,13-ABR-1998 US 60/08957 PR
10-AUG-1998 US 60/096116,04-SEP-1998 US 60/09873 PT LYDIE
10-AUG-1998 US 60/096116,04-SEP-1998 US 60/096116,04-SEP-199
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Bougueleret, L., Duclert, A. and Edwards, J.B.D.M. Elongation cDNA of secretory protein Patent: JP 2001523453-A 47 27-NOV-2001;
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JP 2001523453-A/47
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JP 2001523453-A/47
27-NOV-2001
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Similarity 97.8%;
88; Conservative
                                    TCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
                                                                               GAAAGGCCACGTGA-AGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTAT
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Sig_peptide

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/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 2.2e-103;
4; Mismatches 5;
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7 TGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTC-TGTATTCTT 505	7 TTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAA	7 TGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGC 386	7 TCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGG 326	77 GCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGC 266	8 CGCGATGGATAACGTGCAGCCGAAAATAAAACATC-GCCCCTTCTGCTTCAGTGTGAAAG 206	8 ACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGTTGCAGA 147	8 GGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTCGCAGA 87	h 87.2%; Score 465.8; DB 6; Length 526; Similarity 99.0%; Pred. No. 3.9e-103; 89; Conservative 0; Mismatches 3; Indels 2; Gaps 2;	GGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAG	21 GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC 380

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KEYWORDS
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Local Similarity 99.8%;
nes 412; Conservative
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OS Homo Sapiens (human)

PN JP 2001523453-A/94

PD 27-NOV-2001

PF 13-NOV-1998 JP 2000521191

PF 13-NOV-1998 US 60/066677,17-DEC-1997 US 60/069957 PR

13-NOV-1998 US 60/04121,13-APR-1998 US 60/099273 PI LYDIE

BOUGUELERET, AYMERIC DUCLERT, JEAN BAPTISTE DUMAS MILNE PI EDWARDS

PC C12N15/10,C12N21/09,C07K14/47,C07K16/18,C12N1/15,C12N1/19, PC

C12N1/21,

PC C12N5/10,C12P21/02,C12Q1/68,C12N15/00,C12N15/00 CC

Von Heijne matrix

CC score 4,0999990463257

CC score 4,0999990463257

CC seq VFMLIVSVLALIP/ET

FK KEY

FT sig peptide 46.119

FT polyA_signal 377.382

FT polyA_site 46.119
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Bougueleret, L., Duclert, A. and Edwards, J.B.D.M. Elongation CDNA of secretory protein Patent: JP 2001523453-A 94 27-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BD085952
Elongation cDNA of
BD085952
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Mammalia; Eutheria; Primates;
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                                           GTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGT
                                                                                                   CTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCA
                                                                                                                                                        CTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGCA
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CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAGAAGTTTTGTAATTTTATATTACTTTT 466
                                                                                  CTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCA
                                                                                                                                         CTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCA
                                                                                                                                                                                                CCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCCACGTGAAGATGCTGCGG
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                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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SOURCE
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Edwards, J.B.D.M., Duclair, E. and Jordan,
Sequence tag and encoded human protein
Patent: JP 2001269182-A 138 02-OCT-2001;
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JP 2001269182-A/138
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                          TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                                     CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                                                                                                                                                                                                                                                           GTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
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TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                                           ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
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                                                                                         TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                 CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                                                                                                                                                                                                                                      GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
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Key Local
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26-FEB-1999 US 60/122487
JEAN BAPUTIST DUMAS MILNE EDWARDS, EIMERIC DUCLAIR, JEAN
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02-OCT-2001
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/mo1_type="genomic DNA"
/db_xref="taxon:9606"
97 c 126 q 91
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VERSION
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Homo sapiens chemokine-like f alternatively spliced.
AF136300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
Submitted (17-MAR-1999) Immunology,
Yuan Road, Beijing 100083, China
2 (bases 1 to 689)
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Han, W.L., Gu, W., Li, Y., Zhang, Y., Song, Q.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Han, W.L
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Mammalia; Eutheria; Primates;
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                                                              CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT 194
                                                                                                                                                           GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
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195
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140 c 161 g 193 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/mol_type="mRNA"
/db xref="ftaxon:9606"
/cell_line="U937"
/note="from PHA stimulated
                                                                                                                                                                                                                                                                                                                                                /product="chemokine-like factor 2"
/protein_id="AAF19599.1"
/db_xref="GI:6630854"
                                                                                                                                                                                                                                                                                                                                                                                           /gene="CKLF2"
/note="UCK-2; alternatively spliced"
/codon_start=1
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Pred. No. 1.1e-74;
0; Mismatches 1
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                                                                                                                                                                        Science Center, Xueyuan Road
Sequence update by submitter
On Sep 7, 2000 this sequence
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 593)

Han, W.L., Gu, W.F., Li, Y., Zhang, Y.M., Di, C.H., Song, Q.S. and
                                                                                                                                                                                                                                 Submitted (07-SEP-2000)
                                                                                                                                                                                                                                                                Han, W.L.
                                                                                                                                                                                                                                                                          Direct Submission
Submitted (23-APR-1999) Immunology,
Yuan Road, Beijing 100083, China
2 (bases 1 to 593)
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AF145216.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AF145216 593 bp mRNA linear Homo sapiens chemokine-like factor 4 (CKLF4) mRNA,
                                                                                                                                                                                                                                                Direct Submission
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                                  /gene="CKLF4"
/note="synonym:
148. .510
                                                                                                                      ₽
                                                                                                                                organism="Homo sapiens'
    'note="UCK-4; alternatively spliced"
                   gene="CKLF4"
                                                                                        note="from PHA
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                                                                                                      xref="taxon:9606"
1_line="U937"
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                                                                                                                                                                                                                   Immunology, Peking University Health Road 38, Beijing 100083, China
                                                                                        stimulated
                                                                                                                                                                                          version replaced gi:6625671
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RESULT 10
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Submitted (12-MAR-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                        Homo sapiens, clone MGC:10658
BC004380
BC004380.1 GI:13325133
                                                                              Strausberg, R.
                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                      Homo
                                                            Direct Submission
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YIVITGFEVTVILFFILLYVLRLDRLMKWLFWPLLVFALVTAVCCLADGALIYRKLLF
NPSGFYQKKPVHEKKEVL"
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/product="chemokine-like factor
/protein_id="AAF19350.1"
/db_xref="GI:6625672"
                                                                                             to 688)
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Pred. No. 1.3e-74;
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8 IMAGE:3639550, mRNA,
                                                                                                                Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 9989692.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiù, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Marrin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prahbu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Genome Sequence Centre,
RC Cancer Agency, Vancouver, BC, Canada
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Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         George Yang,
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  -GATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCACT
                                          CATACTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCT
                                                                                                                        ACAAGCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTT
                                                                                                                                                                                                         CCACGTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGC
                                                                                                                                                                                                                                               CCACGTGAAGATGCTGCGGCTG-----
                                                                                                                                                                                                                                                                                         CGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGG
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/product="Unknown (protein protein_id="AAH04380.1"
/b_xref="GI:13325134"
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|mol_type="mRNA"
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clone_lib="NIH_MGC_39"
lab_host="DH10B-R"_
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Best Local Similarity 75.6
Matches 495; Conservative
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1 (bases 1 to 655)
Zhang,J.S., Nelson,M., Wang,L. and Smith D.I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                   CCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCT
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 TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGGT
                          CCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCCGCCGCGGTGGCGGTTGCTATCGCT
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137 c 153 g 194 t
                                                                                                                                                                                                                                                       /note="down regulated upon sodium butyrate"
                                                                                                                                                                                                                                                                                               /gene="C32"
132. .590
                                                                                                                                                                                                 /product="transmembrane proteolipid"
/protein_id="AAF21255.1"
/db_xref="GI:6648619"
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                                                                                                                                                                                                                                                                                gene="C32"
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                                                                                                                                                                                                                                                                                                                                                 tissue_type="pancreas"
                                                                                                                                                                                                                                                                                                                                                              cell_line="AsPC-1"
                                                                                                                                                                                                                                                                                                                                                                                                     organism="Homo sapiens"
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db_xref="taxon:9606"
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Edwards, J.-B.D.M., Bougueleret, L. and Jobert, S.
Complementary DNA's encoding proteins with signal
Patent: US 6548633-A 70 15-APR-2003;
                                                                                                                                                                                                                                                                                                          Unclassified.
                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                         Unknown
                                                                                                                                                                                                                                                                                                                                     Unknown
                                                                                             GTGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
 CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
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Pred. No. 5.9e-68;
0; Mismatches 2;
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PAT 12-JUN-2003

655

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161;

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                                                                                                                                                                                                                                                                                                                                                                Sequence 70 from AX061665
                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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  /note="Von . 630 . . 635 655 . . 669
                                                 /proteIn_id="CAC25025.1"
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/db_xref="GI:12406790"
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YIVITGEBVTVILFFILLXYVLRLDRLKKMLFWPLLDIINSLVTTVFMLIVSVLALIPE
TTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGFYQKKPVHEKKEVL"
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                                                                                                                                                   /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 434)

Han,W.L., Gu,W., Li,Y., Zhang,Y., Song,Q., Di,C. and Ma,D. Direct Submission
Submitted (17-MAR-1999) Immunology, Beijing Medical University, Yuan Road, Beijing 100083, China
                                                                                                                                                  434 bp mRNA linear Homo sapiens chemokine-like factor 3 (CKLF3) mRNA, alternatively spliced.
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                                                                                        Homo sapiens (human)
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Han,W.L.
Direct Submission
Submitted (07-SEP-2000) Immunology, Peking University Health
Science Center, Xueyuan Road 38, Beijing 100083, China
Sequence update by submitter
On Sep 7, 2000 this sequence version replaced gi:6630855.
Location/Qualifiers
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  G30204
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                                                                                      AAAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTTTGATACTAAGTATTAAACATATT
                                                                                                                                                                           TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/cell_line="U937"
/note="from PHA stimulated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /codon_start=1
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/protein_d="AAF19600.1"
/db_xref="G1:6630856"
/translation="MDNVQPKIKHRPFCFSVKGHVKMLRLVFALVTAVCCLADGALIY
RKLLFNPSGPYQKKPVHEKKEVL"
a 88 c 116 g 103 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /gene="CKLF3"
/note="UCK-3; alternatively spliced"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene="CKLF3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note="UCK"
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Pred. No. 1.1e-66;
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  STS 05-OCT-1996
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Best Local Similarity
Matches 281; Conserv
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primer_bind
primer_bind
E COUNT 1
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Fax: 4157259689
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                                                                                                                                                                                                               146
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                                                                                                                                                    Conservative
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KCl:
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DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
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Mammalia; Eutheria; I
1 (bases 1 to 485)
Myers,R.M.
Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human STS
G30204
G30204.1
STS; STS 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer A: ACTTCTTTTTTTCATGCACAGG
Primer B: GCCTTATTTACCGGAAGCT
STS size: 77
PCR Profile:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Richard M. Myers
Stanford Human Genome Center (SHGC)
Stanford University School of Medicine
Department of Genetics, M-344, Stanford,
                                                                                                                                                                                                                                                                                                                                            Prepared with primer pairs provided by Sandoz,
-- Washington University/Merck EST sequence.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: myers@shgc.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
             AGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGG
                                                                ACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGC
                                                                                                                     GCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGC
AGTATGCTGTCTTGCCGACGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGG
                                                   ACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGC
                                                                                                       GCTTGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTANCTGTGTTGGC
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Taq Polymerase:
Total Vol:
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PCR Cycles:
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                                                                                                                                                                                                                     complement(126. .145)
1 99 c 99 g
                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Primates;
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Catarrhini; Hominidae; Homo
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Sear Job 1	망	Ş	DЬ	γQ
Search completed: January 31, 2004, 03:34:57 Job time : 2120 secs	47 TTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTATAATAAA 1	466 TTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCAAAAA 512	107 TCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTACTTT 48	406 TCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAGAAGTTTTGTAATTTATATTACTTT 465

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Result
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
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BG
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seq length: 2000000000
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Gapop 10.0 , Gapext 1.0
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        94.9
93.3
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SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1987.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1987.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1989.DAT: *

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SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1995.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1995.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1997.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT: *

SIDS1/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT: *
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Copyright (c) 1993 - 2004 Compugen Ltd.
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/SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT:*
/SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1983.DAT:*
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/SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:*
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                                                                                                                                                                                                                                    Length
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AAS44932
AAH98515
AAH98548
AAH34835
AAAH37730
AAF64012
        AAV59598
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Human colon cancer
Human secreted pro
cDNA encoding huma
Human secreted pro
                                                                                                   UCK-1 nucleotide s
cDNA encoding nove
Human EST-derived
Human EST-derived
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Human	EST	AAC78499	21	495	34.7	185.2		
9 486.6 91.1 504 24 ABS73585 11 478.6 89.6 506 12 AAX79746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ARX36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAC00147 17 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 22 AAX9493 20 335 62.7 663 22 AAX9493 21 327.2 61.3 908 22 AAX9493 22 321.8 60.3 669 21 AAA64933 23 321.8 60.3 669 22 AAA64933 24 309.8 58.0 354 20 AAX41509 25 300 55.2 566 22 AAX41509 26 305.6 57.4 670 24 AAB15919 27 305.8 57.3 638 24 ABX35884 28 300 56.2 566 22 AAX45120 29 300 56.2 566 22 AAX45120 20 33 224.4 42.0 459 21 AAA38007 31 224.4 40.1 13744 22 AAX682407 34 214.4 40.1 13744 22 AAX68361 39 205 38.4 207 24 ABL62782 40 205 38.4 207 24 ABL62782 41 205 38.4 207 24 ABL62782 42 205 38.4 207 24 ABL62782 43 201.4 37.7 326 24 ABL63783 45 201.4 37.7 326 24 ABL63783 46 201.4 37.7 326 24 ABL63783 47 201.4 37.7 326 24 ABL63783 48 201.4 37.7 326 24 ABL63783 49 205 38.4 207 24 ABL63783 40 201.4 37.7 326 24 ABL63783 41 201.4 37.7 326 24 ABL63783 42 201.4 37.7 326 24 ABL63783 43 201.4 37.7 326 24 ABL63783 44 201.4 37.7 326 24 ABL63783 45 201.4 37.7 326 24 ABL63783 46 201.4 37.7 326 24 ABL63783 47 201.4 37.7 326 24 ABL63783 48 201.4 37.7 326 24 ABL63783 49 205 38.4 207 24 ABL63783 40 201.4 37.7 326 24 ABL63783 41 201.4 37.7 326 24 ABL63783 42 201.4 37.7 326 24 ABL63783 43 201.4 37.7 326 24 ABL63783 44 201.4 37.7 326 24 ABL63783 45 201.4 37.7 326 24 ABL63783 46 201.4 37.7 326 24 ABL63783 47 201.4 37.7 326 24 ABL63783 48 201.4 37.7 326 24 ABL63783 49 201.4 37.7 326 24 ABL63783 40 201.4 37.7 326 24 ABL63783 41 201.4 37.7 37.8 ABL63783 41	Human EST DNA435	AAZ34052	20	495	ω 4.	185.2		
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 12 AAV59746 Human secretee 11 478.6 89.6 506 22 AAX57826 Human cDNA #2 14 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ABX36006 Human secretee 15 411.4 77.0 413 20 AAX97873 Human secretee 16 405.4 75.9 415 21 AAC0147 Human secretee 17 335 62.7 663 21 AAC78498 18 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 20 335 62.7 663 21 AAC78498 21 327.2 61.3 908 22 AAX9451 22 321.8 60.3 669 22 AAX94933 Human PRO772 1 23 321.8 60.3 669 22 AAX64933 Human secretee 25 30.0 56.5 57.4 670 24 ABL90366 CDNA encoding 26 306.6 57.4 670 24 ABL90366 CDNA encoding 27 305.8 57.3 638 24 ABX35884 CDNA encoding 28 300 56.2 566 22 AAX45120 CDNA encoding 29 300 56.2 566 22 AAX45120 CDNA encoding 30 295.4 55.3 297 21 AAA15919 Human protein 31 285 53.4 207 AAX68361 Human protein 32 221.4 40.1 13744 22 AAX68361 Human protein 33 221.4 40.1 13744 22 AAX68361 Human immune/I Human immune/I Human secretee 40 205 38.4 207 24 ABL62782 Human protein Human immune/I Human immune/I Human immune/I Human immune/I Human immune/I Human immune/I Human secretee Human protein Human protein Human protein Human protein Human protein Human protein		713	24	326	37.	201.4		
9 486.6 91.1 504 24 ABS73585 11 478.6 89.6 506 19 ABS73739 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ABX36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAA234051 17 335 62.7 663 21 AAA234051 18 335 62.7 663 22 AAX9423 21 327.2 61.3 908 22 AAX9423 22 321.8 60.3 669 21 AAA24933 231.8 60.3 669 21 AAA24053 24 309.8 58.0 354 20 AAX91509 25 30.6 57.4 67.0 21 AAA254053 26 30.8 57.3 638 24 ABX36046 27 305.8 57.3 638 24 ABX35046 28 300 56.2 566 22 AAA34933 30 256.2 566 22 AAA34933 31 285 53.4 544 23 ABV56390 32 24.4 42.0 45.9 12 AAA35919 33 27.4 42.0 45.9 12 AAA36907 34 214.4 40.1 13744 22 AAX68361 36 210.2 39.4 20.7 24 ABL62959 37 20 38.4 20.7 24 ABL62959 38 29 38.4 20.7 24 ABL62959 38 29 38.4 20.7 24 ABL62959 38 29 38 29 38 38 38 38 38 38 38 38 38 38 38 38 38	cancer	318	24	207	38.	205		ი
9 486.6 91.1 504 24 ABS73585 11 478.6 89.6 506 12 AAX59746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ABX36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAC00147 17 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 22 AAX9493 20 335 62.7 663 21 AAC78498 21 327.2 61.3 908 22 AAX64933 21 327.2 61.3 908 22 AAX64933 21 321.8 60.3 669 22 AAX64933 22 331.8 60.3 669 22 AAX64933 23 321.8 60.3 669 22 AAX64053 24 309.8 58.0 354 20 AAX41509 25 300 55.2 566 22 AAX645120 26 300 56.2 566 22 AAX65121 27 305.8 57.3 638 24 ABL90366 28 300 56.2 566 22 AAX65121 29 300 56.2 566 22 AAX65120 20 33 224.4 42.0 459 21 AAA38007 34 214.4 40.1 13744 22 AAX68361 36 214.4 40.1 13744 22 AAX68361 37 214 40.1 13744 22 AAX68361 38 205 38.4 207 24 ABL907651 39 205 38.4 207 24 ABL907651 30 205 38.4 207 24 ABL60782 30 205 38.4 207 24 A	cancer	ABL62959	24	207	38.	205		ი
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 22 AAX57746 Human secretec 11 478.6 89.6 506 24 ABS73739 Human cDNA #2 12 466.8 87.4 500 20 AAX97826 Human secretec 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ABX36006 Human secretec 15 411.4 77.0 413 20 AAX97873 Human secretec 16 405.4 77.0 413 20 AAX97873 Human secretec 17 335 62.7 663 21 AAC78498 Human secretec 18 335 62.7 663 21 AAC78498 Human proote in 19 335 62.7 663 21 AAC78498 Human proote in 20 335 62.7 663 22 AAX94933 Human proote in 21 327.2 61.3 908 22 AAX94933 Human proote in 22 321.8 60.3 669 22 AAX94933 Human secretec 23 321.8 60.3 669 22 AAX94933 Human secretec 24 309.8 58.0 354 20 AAX41509 Human secretec 25 30.3 669 22 AAX96493 Human secretec 26 306.6 57.4 670 24 ABL90366 Human secretec 27 305.8 57.3 638 24 ABX35884 Human secretec 28 300 56.2 566 22 AAX96493 Human secretec 29 300 56.2 566 22 AAX96493 Human secretec 20 301 52.2 566 22 AAX96493 Human secretec 20 302 52.4 670 24 ABL90366 Human prootein 20 303 56.2 566 22 AAX95120 Human prootein 21 AAX56930 Human prootein 22 AAX69361 Human prootein 23 221.4 42.0 621 22 AAX68361 Human prootein 24 40.1 13744 22 AAX68361 Human immune/! Human immune/! Human secretec 25 38.4 207 24 ABN39666 Human immune/! Human immune/! Human secretec 26 27 AAX97663 Human immune/! Human immune/! Human secretec 27 AAX97663 Human immune/! Human secretec 28 AAX97663 Human immune/! Human immune/! Human secretec Extended human	ncer	ABL62782	24	207	38.	205		Ω
9 486.6 91.1 504 24 ABS73585 11 478.6 89.6 506 19 ABA79746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 ABA15929 14 425 79.6 43 24 ABX36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAC78498 17 335 62.7 663 21 AAC78498 18 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 22 AAX9423 21 327.2 61.3 908 22 AAX9423 22 321.8 60.3 669 21 AAC84053 23 321.8 60.3 669 21 AAA87771 24 309.8 58.0 354 20 AAX41509 25 309 57.9 637 21 AAC84053 26 306.6 57.4 670 24 ABL90366 27 305.8 57.3 638 24 ABA59121 29 300 56.2 566 22 AAS45121 30 30 56.2 566 22 AAS45121 31 285 53.4 544 23 ABX56007 31 285 53.4 544 23 ABX56007 31 284 40.1 13744 22 AAX68361 31 1374 21 40.1 13744 22 AAX68360 25 AAX60850 36 21 AAC68850 26 Extended human immune// 37 214 40.1 269 21 AAX66850 27 Extended human immune// 38 210 25 24 24 AAX60850 27 Extended human immune// 38 210 25 24 AAX60850 28 21 AAX60850 28 Extended human immune// 39 210.2 39.4 321 20 AAX97651 28 21 21 21 AAX16800 28 21 AAX608406 29 21 AAX608406 20 21 AAX608406 21 AAX608406 21 AAX608406 21 AAX608406 21 AAX608406 22 AAX97651 23 39.4 321 20 AAX97651 28 28 39.4 32.1 20 AAX97651	used	ABN94246	24	207	38.	205		Ω
9 486.6 91.1 504 24 ABS73585 10 478.6 89.6 506 19 AAV59746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ARX36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAC0147 17 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 22 AAX97873 20 335 62.7 663 21 AAC78498 21 327.2 61.3 908 22 AAX9423 22 321.8 60.3 669 22 AAX9423 23 321.8 60.3 669 22 AAX96405 24 309.8 58.0 354 20 AAX41509 25 309.8 58.0 354 20 AAX41509 26 300.5 57.4 670 24 AAEX56747 27 305.8 57.3 638 24 ABX35884 28 300 56.2 566 22 AAX5121 29 300 56.2 566 22 AAX5121 30 295.4 55.3 297 21 AAX15919 31 285 53.4 939 22 AAX85210 32 224.4 42.0 459 21 AAX80407 34 214.4 40.1 13744 22 AAX80361 37 214 40.1 13744 22 AAX8046 37 214 40.1 13744 22 AAX8026 38 21 AAX008850 39 22 AAX8026 30 22 AAX802	Extended human s	AAX97651	20	321	39.	210.2		
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 21 AAV59746 Human secretee 11 478.6 89.6 506 22 AAX597826 Human cDNA #2 12 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 Human secretee 14 425 79.6 439 22 AAX34051 Human secretee 15 411.4 77.0 413 20 AAX97873 Human secretee 16 405.4 77.0 413 20 AAX397873 Human secretee 17 335 62.7 663 21 AAC78498 Human proof a Human proof a Formal Proof		AAC06850	21	269	40.	214		
9 486.6 91.1 504 24 ABS73585 11 478.6 89.6 506 19 ABA79746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 43 24 ABX36006 15 411.4 77.0 413 20 AAX37873 16 405.4 75.9 415 21 AAA234051 17 335 62.7 663 21 AAA234051 18 335 62.7 663 21 AAA27423 21 327.2 61.3 908 22 AAX4423 22 321.8 60.3 669 21 AAA84933 21 327.2 61.3 908 22 AAX4973 23 321.8 60.3 669 21 AAA87771 24 309.8 58.0 354 20 AAX41509 25 309.8 57.9 637 21 AAA56747 26 306.6 57.4 670 24 ABL50366 27 305.8 57.3 638 24 ABX35884 28 300 56.2 566 22 AAS45121 39 39 57.9 637 21 AAA55120 29 30 56.2 566 22 AAS45121 30 295.4 55.3 297 21 AAA16090 31 285 53.4 544 23 ABX95630 32 24.4 40.1 13744 22 AAK68361 Human prootacte 13 405.4 40.1 13744 22 AAK68361 Human colon cc 14 AB.6 89.6 91 AAA8930 14 AAA668361 Human colon cc 15 ABA668361 Human colon cc 16 ABA76764 17 ABA6684053 Human colon cc 17 ABA6684053 Human colon cc 18 ABA76764053 Human colon cc 18 ABA76764054 Human colon cc 18 ABA76764054 Human colon cc 18 ABA76764054 Human colon cc 18 ABA76764 Human immune//		AAK82406	22	13744		214.4	36	
9 486.6 91.1 504 24 ABS73585 10 478.6 89.6 506 19 AAV59746 11 478.6 89.6 506 24 ABS73739 12 466.8 87.4 500 20 AAX97826 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 ARA36006 15 411.4 77.0 413 20 AAX97873 16 405.4 75.9 415 21 AAC00147 17 335 62.7 663 21 AAC78498 18 335 62.7 663 21 AAC78498 19 335 62.7 663 22 AAX97873 18 335 62.7 663 22 AAX97873 20 335 62.7 663 22 AAX97873 21 327.2 61.3 908 22 AAX9423 22 321.8 60.3 669 21 AAC78493 23 321.8 60.3 669 21 AAA678493 24 309.8 58.0 354 20 AAX41509 25 305.6 57.4 670 24 ABL90366 27 305.8 57.3 638 24 AAB35804 29 300 56.2 566 22 AAS45121 30 295.4 55.3 297 21 AAA15919 31 285 53.4 544 23 ABV56930 31 284.4 42.0 459 21 AAA38007 31 21.4 6021 22 AAX682407 31 AAA16090 31 21.4 4 40.1 6021 22 AAX682407 31 AAA16090 31 21.4 40.1 6021 22 AAX682407	immune/	AAK68361	22	13744	•	214.4	35	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 21 AAV59746 Human secretee 11 478.6 89.6 506 22 AAX97826 Human cDNA #2 12 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 14 425 79.6 439 24 AAX97873 16 405.4 77.0 413 20 AAX97873 17 411.4 77.0 413 20 AAX97873 18 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 19 335 62.7 663 21 AAC78498 20 335 62.7 663 21 AAC78498 21 327.2 61.3 908 22 AAS44933 22 321.8 60.3 669 22 AAS4493 23 321.8 60.3 669 22 AAF64053 24 309.8 58.0 354 20 AAX41509 25 300 56.2 566 22 AAX41509 26 306.6 57.4 670 24 ABL90366 27 305.8 57.3 638 24 ABX35884 28 300 56.2 566 22 AAS45120 29 300 56.2 566 22 AAS45120 29 300 56.2 566 22 AAS45120 29 300 56.2 566 22 AAS45120 20 30 55.3 59.4 59.3 ABV56930 31 285 53.4 42.0 459 21 AAA38007 UCK-2 mucleotit	_	AAK82407	22	6021	40.1	214.4	34	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 24 ABS73739 Human secretecy 11 478.6 89.6 506 22 AAX97826 Human secretecy 12 466.8 87.4 500 20 AAX97826 Human secretecy 13 465.4 87.2 467 21 AAA15929 Human secretecy 14 425 79.6 43 24 ABX36066 Human protein 15 411.4 77.0 413 20 AAX97873 Human secretecy 15 411.4 77.0 413 20 AAX97873 Human secretecy 16 405.4 75.9 415 21 AAC0147 Human secretecy 17 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC78493 CDNA encoding 24 309.8 58.0 354 20 AAX41509 Human pro772 r 23 321.8 60.3 669 22 AAF64053 ABX94933 CDNA encoding 24 309.8 58.0 354 20 AAX41509 Human secretecy 25 309 57.4 670 24 ABL59366 Human secretecy 26 300.5 57.4 670 24 ABL59366 Human propositatesy 27 305.8 57.3 638 24 ABA55120 Human propositatesy 28 300 56.2 566 22 AAS45121 Human propositatesy 29 300 56.2 566 22 AAS45090 Human propositatesy 20 302 272 50.9 392 21 AAA16090 Human propositatesy Human propo		AAA38007	21	459	42.0	224.4	33	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 12 AAV57379 Human secretee 11 478.6 89.6 506 24 ABS73739 Human secretee 12 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 Human secretee 14 425 79.6 43 24 ABX36006 Human secretee 15 411.4 77.0 413 20 AAX97873 Human secretee 16 405.4 75.9 415 21 AAC00147 Human secretee 17 335 62.7 663 21 AAC78498 Human secretee 18 335 62.7 663 21 AAC78498 Human pRO772 r 18 335 62.7 663 21 AAC78498 Human pRO772 r 19 335 62.7 663 22 AAX94051 Human pRO772 r 19 335 62.7 663 25 AAX9423 CDNA encoding 20 335 62.7 663 25 AAX9453 21 327.2 61.3 908 22 AAX9453 22 321.8 60.3 669 21 AAA39771 Human pro772 r 23 321.8 60.3 669 21 AAA39771 Human secretee 25 309.8 58.0 354 20 AAX91509 Human secretee 26 306.6 57.4 670 24 ABL90366 Human secretee 27 305.8 57.3 638 24 ABS3584 Human polynucly 28 300 56.2 566 22 AAS45120 CDNA encoding 29 300 56.2 566 22 AAS45121 CDNA encoding 29 300 56.2 566 22 AAS45121 CDNA encoding 20 20 30.4 55.3 297 21 AAA15919 Human protein 31 285 53.4 544 23 ABV56930 Human protein	_	AAA16090	21	392	50.9	272	32	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 21 AAV59746 Human secretee 11 478.6 89.6 506 22 AAX57826 Human secretee 12 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 Human secretee 14 425 79.6 43 24 AAX97873 Human secretee 15 411.4 77.0 413 20 AAX97873 Human secretee 16 405.4 75.9 415 21 AAC00147 Human secretee 17 335 62.7 663 21 AAC78498 Human PRO772 r 18 335 62.7 663 21 AAC78498 Human PRO772 r 19 335 62.7 663 21 AAC78498 Human PRO772 r 20 335 62.7 663 21 AAX97873 Human PRO772 r 21 327.2 61.3 669 22 AAS44933 CDNA encoding 22 321.8 60.3 669 22 AAF64053 Human secretee 23 321.8 60.3 669 22 AAF64053 CDNA encoding 24 309.8 58.0 354 20 AAX41509 Human secretee 25 309.8 57.9 637 21 AAE56747 Human secretee 26 306.6 57.4 670 24 AAE56747 Human secretee 27 305.8 57.3 638 24 ABK35884 CDNA encoding 28 300 56.2 566 22 AAS45120 CDNA encoding 29 300 56.2 566 22 AAS45120 CDNA encoding 20 305.4 55.3 297 21 AAA545919 Human pro772 r 20 AAS45120 CDNA encoding 20 300 56.2 566 22 AAS455120 CDNA encoding 21 AAA545121 CDNA encoding 22 AAS45120 CDNA encoding 23 CDNA encoding 24 AAS45120 CDNA encoding 25 AAS45120 CDNA encoding 26 CDNA encoding 27 AAA545120 CDNA encoding 28 AAA455120 CDNA encoding 29 300 56.2 566 22 AAS455120 CDNA encoding 29 300 56.2 566 22 AAS455120 CDNA encoding 29 300 56.2 566 22 AAS455120 CDNA encoding		ABV56930	23	544	53.4	285	31	ი
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 24 ABS73739 Human secretec 11 478.6 89.6 506 24 ABS73739 Human secretec 12 466.8 87.4 500 20 AAX97826 Human secretec 13 465.4 87.2 467 21 AAA15929 Human secretec 14 425 79.6 439 24 ABA3606 Human secretec 15 411.4 77.0 413 20 AAX97873 Human secretec 16 405.4 75.9 415 21 AAC0147 Human secretec 17 335 62.7 663 20 AAX34051 Human secretec 18 335 62.7 663 21 AAC78498 Human pRO772 r 18 335 62.7 663 21 AAC78498 Human pRO772 r 18 335 62.7 663 21 AAC82423 CDNA encoding 21 327.2 61.3 908 22 AAX41593 CDNA encoding 22 321.8 60.3 669 21 AAX41509 AAX41509 AAX41509 Human secretec 23 301.8 60.3 669 21 AAX41509 Human secretec 24 309.8 58.0 354 20 AAX41509 Human secretec 25 309.8 58.0 354 20 AAX41509 Human secretec 26 306.6 57.4 670 24 ABL90366 CDNA encoding 27 305.8 57.3 638 24 ABX55884 Human polynucl 28 300 56.2 566 22 AAS45121 CDNA encoding	protein	AAA15919	21	297	55.3	295.4	30	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 24 AAS73739 Human secretecel 11 478.6 89.6 506 22 AAX97826 Human secretecel 12 466.8 87.4 500 20 AAX97826 Human secretecel 13 465.4 87.2 467 21 AAA15929 Human secretecel 14 425 79.6 439 24 ABX36006 Human protein 14 425 79.6 439 24 AAX37873 Human secretecel 15 411.4 77.0 413 20 AAX37873 Human secretecel 16 405.4 75.9 415 21 AAC78498 Human secretecel 17 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC78498 Human pro772 r 19 335 62.7 663 21 AAC78498 Human pro772 r 20 335 62.7 663 21 AAC78493 Human pro772 r 21 327.2 61.3 908 22 AAX4453 J321.8 60.3 669 21 AAA87771 J331.8 60.3 669 21 AAA87771 J341.8 60.3 669 21 AAA87771 J352.3 321.8 60.3 669 21 AAA87771 J373.3 58.0 354 20 AAX41509 Human secretecel 25 30.0 57.9 637 21 AA256747 Human pro72 r 26 305.8 57.3 638 24 ABL50366 Human pro72 r 27 305.8 57.3 638 24 ABL50366 Human proplymud 28 300 56.2 566 22 AAS45120 CDNA encoding	encoding	AAS45121	22	566	56.2	300	29	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 12 AAX59746 Human secretee 11 478.6 89.6 506 24 AAX597826 Human secretee 12 466.8 87.4 500 20 AAX97826 Human secretee 13 465.4 87.2 467 21 AAA15929 Human secretee 14 425 79.6 43 24 AAX37873 Human secretee 15 411.4 77.0 413 20 AAX97873 Human secretee 16 405.4 75.9 415 21 AAC00147 Human secretee 17 335 62.7 663 21 AAC78498 Human PRO772 1 18 335 62.7 663 21 AAC78498 Human PRO772 1 19 335 62.7 663 21 AAC78498 Human PRO772 1 20 335 62.7 663 21 AAX97873 Human PRO772 1 21 327.2 61.3 908 22 AAX9493 22 321.8 60.3 669 21 AAX9493 23 321.8 60.3 669 21 AAA6493 24 309.8 58.0 354 20 AAX41509 Human secretee 25 306.6 57.4 670 24 ABS35884 CDNA encoding 26 306.6 57.4 670 24 ABS35884 CDNA sequence	encoding	AAS45120	22	566	56.2	300	28	
9 486.6 91.1 504 24 ABS73585 Human cDNA #1 10 478.6 89.6 506 19 AAV59746 Human secretec 11 478.6 89.6 506 24 ABS73739 Human secretec 12 466.8 87.4 500 20 AAX97826 Human secretec 13 465.4 87.2 467 21 AAA15929 Human secretec 14 425 79.6 439 24 ABS46006 Human protein 15 411.4 77.0 413 20 AAX97873 Human secretec 15 411.4 77.0 413 20 AAX97873 Human secretec 16 405.4 75.9 415 21 AAC00147 Human secretec 17 335 62.7 663 20 AA234051 Human pro772 r 18 335 62.7 663 21 AAC78498 Human pro772 r 18 335 62.7 663 21 AAC58238 Human pro772 r 20 321.8 60.3 669 22 AAA84771 CDNA encoding 21 327.2 61.3 908 22 AAA84771 Human secretece 23 321.8 60.3 669 22 AAF64053 Human secretece 24 309.8 58.0 354 20 AAX35674 Human secretece 25 309 57.9 637 21 AA256747 Human pro772 r 26 306.6 57.4 670 24 ABL90366 Human pro772 r 27 ABL90366 Human secretece 28 306.6 57.4 670 24 ABL90366 Human prodynucl	sequence	ABK35884	24	638	57.3	305.8	27	
486.6 91.1 504 24 ABS73585 478.6 89.6 506 19 AAV59746 478.6 89.6 506 24 ABS73739 466.8 87.4 500 20 AAX97826 465.4 87.2 467 21 AAA15929 425 79.6 439 24 ABS73606 411.4 77.0 413 20 AAX97873 405.4 75.9 415 21 AAC00147 405.4 75.9 415 21 AAC00147 405.4 75.9 415 21 AAC34051 335 62.7 663 20 AAX294031 335 62.7 663 21 AAC78428 336 62.7 663 22 AAS44933 327.2 61.3 908 22 AAS4933 327.2 61.3 908 22 AAS4933 321.8 60.3 669 22 AAS4933 321.8 60.3 669 22 AAS65747 Human secretee CDNA encoding Human secretee Human secretee CDNA encoding Human secretee		ABL90366	24	670	57.4	306.6	26	Ω
486.6 91.1 504 24 ABS73585 478.6 89.6 506 19 AAV59746 478.6 89.6 506 24 ABS73739 466.8 87.4 500 20 AAX97826 465.4 87.2 467 21 AAAX5929 411.4 77.0 413 20 AAX97873 411.4 77.0 413 20 AAX97873 405.4 75.9 415 21 AAC00147 405.5 47.5 663 20 AAX24051 335 62.7 663 21 AAC78498 335 62.7 663 21 AAC78498 335 62.7 663 21 AAC58238 335 62.7 663 21 AAC58243 337.2 61.3 908 22 AAS4493 321.8 60.3 669 22 AAS44771 321.8 60.3 669 22 AAX41509 408.4 89.6 500 4000 4000 4000 4000 4000 4000 4000		AAZ56747	21	637	57.9	309	25	
486.6 91.1 504 24 ABS73585 Human cDNA #1 478.6 89.6 506 19 AAV59746 Human secretee 478.6 89.6 506 24 ABS73739 Human cDNA #2 466.8 87.4 500 20 AAX57826 Human secretee 465.4 87.2 467 21 AAA15929 Human protein 425 79.6 439 24 ABK56006 CDNA sequence 411.4 77.0 413 20 AAX97873 Human secretee 411.4 77.0 413 20 AAX97873 Human secretee 411.4 77.0 663 21 AAC0147 Human protein 335 62.7 663 21 AAC78498 Human pro772 r CDNA encoding 321.8 60.3 669 21 AA85493 321.8 60.3 669 22 AAF64053 CDNA encoding		AAX41509	20	354	58.0	309.8	24	
486.6 91.1 504 24 ABS73585 478.6 89.6 506 19 AAV59746 478.6 89.6 506 24 ABS73739 466.8 87.4 500 20 AAX97826 465.4 87.2 467 21 AAA15929 425 79.6 439 24 ABX36006 411.4 77.0 413 20 AAX97873 405.4 75.9 415 21 AAC00147 335 62.7 663 20 AAZ94051 335 62.7 663 21 AAC58238 337.2 61.3 908 22 AAA487771 408.6 91.1 504 248873771 408.7 4	Ф	AAF64053	22	669	60.3	321.8	23	
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486.6 91.1 504 24 ABS73585 478.6 89.6 506 19 AAV59746 478.6 89.6 506 24 ABS73739 466.8 87.4 500 20 AAX97826 465.4 87.2 467 21 AAA15929 425 79.6 439 24 ABX36006 411.4 77.0 413 20 AAX97873 Human cDNA #2 AAX97873 Human protein	secrete	AAC00147	21	415	•	405.4	16	
486.6 91.1 504 24 ABS73585 Human cDNA #1 478.6 89.6 506 19 AAV59746 Human secretece 478.6 89.6 506 24 ABS73739 Human cDNA #2 466.8 87.4 500 20 AAX97826 Human secretece 465.4 87.2 467 21 AAA15929 Human protein 425 79.6 439 24 ABK36006 cDNA sequence	secreted	AAX97873	20	413	•	411.4	15	
486.6 91.1 504 24 ABS73585 Human cDNA #1 478.6 89.6 506 19 AAV59746 Human secretee	sequence	600	24	439	٠	N	14	
486.6 91.1 504 24 ABS73585 Human cDNA #1 478.6 89.6 506 19 AAV59746 Human cDNA #2 478.6 89.6 506 24 ABS73739 Human cDNA #2 466.8 87.4 500 20 AAX57826 Human secretce	protein	N	21	467	•	65.	13	
486.6 91.1 504 24 ABS73585 Human cDNA #1 478.6 89.6 506 19 AAV59746 Human secreted 478.6 89.6 506 24 ABS73739 Human cDNA #2	secreted	N	20	500	•	66.	12	
486.6 91.1 504 24 ABS73585 Human cDNA #1 f 478.6 89.6 506 19 AAV59746 Human secreted	cDNA #2	73	24	506	•	78.	11	
486.6 91.1 504 24 ABS73585 Human cDNA #1 f	secreted	74	19	506	•	78.	10	
	CDNA #1 f	735	24	504	•	86.	φ	

ALIGNMENTS

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WPI; 2000-388170/34.
P-PSDB; AAY98142.
                                  Ma D,
                                                                                                                                 UCK-1; chemotaxis factor; immunocyte; haemopoiesis stimulant; tumour; radiotherapy; chemotherapy; human; ss.
Chemotaxic factor useful for treatment and diagnosis of immunocyte
                                                              14-MAY-1999;
                                                                            14-MAY-1999;
                                                                                          16-FEB-2000
                                                                                                        CN1244584-A
                                                                                                                     Homo sapiens
                                                (UYBE-) UNIV BEIJING MEDICAL
                                  Han W,
                                 Zhang
                                                              99CN-0107284.
                                                                            99CN-0107284.
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UCK-1 nucleotide sequence.

22-AUG-2000 (first entry)

AAA38006;

AAA38006 standard; cDNA; 534

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Best Local S
Matches 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents an UCK-1 cDNA sequence encoding a chemotaxis factor polypeptide. The UCK-1 protein exhibits immunocyte chemotaxis activity and a haemopoiesis stimulating effect. The invention relates to UCK proteins, their encoding nucleotide sequences and antibodies and antagonists against the proteins. The nucleotide and protein sequences are useful for the preparation of a composition for the diagnosis and treatment of diseases associated with abnormal immunocyte function and low haemopoiesis function caused by radiotherapy and chemotherapy used
transgenic animal; Alzheimer's disease; Parkinson's disease; burn; amyotrophic lateral sclerosis; platelet disorder; thrombocytopenia; ulcer; osteoporosis; bone degenerative disorder; periodontal disease; gut protection; lung; liver fibrosis; immune deficiency; infection;
                                                                                              cDNA encoding
                                                                                                                                                                               AAS44932 standard; cDNA; 655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 534 BP; 156 A; 110 C;
                                                      ischaemia-reperfusion injury; haematopoiesis;
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                                                                                                                                                                                                                                                                     GCCTGTGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTTGATACTAA
                                                                                                                                                                                                                                                                                                                                                                                CGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                               CACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGC
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                                                                                                                                                                                                                                                                                                                                                                                                                    CACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGC
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                                                                                                                          (first
                                                                                              novel human secretory
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                                                                   protein; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunocyte chemotaxic stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31pp; Chinese.
                                                                                                                         entry)
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Pred. No. 2.3e-107;
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                                                                   Crohn's
                                                                                            protein,
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                                                      cancer; neuropathy;
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The invention relates to novel isolated human secreted polypeptides (I) CC and polynucleotides (II). (I) and (II) are useful for treating CC inflammatory conditions such as arthritis, nephritis, Crohn's disease, CC ischaemia-reperfusion injury, shock, sepsis, immune responses, and is CC involved in increasing haematopoiesis, stem cell survival, bone growth CC and remodeling. (I), (II) and modulators of (II) are useful for CC prophylaxis or treatment of one or more cancers. (II) is also useful for CC creating transgenic animals useful for studying the in vivo activities of the polypeptide as well as for studying modulators of the polypeptides. (CC II) induces the proliferation of neural cells and regeneration of nerve and brain tissue and is useful for the treatment of central and CC parkinson's disease, Huntington's disease, and amyotrophic lateral CC parkinson's disease, Huntington's disease, and amyotrophic lateral CC cativity, regulation of haematopoiesis and is useful for treating myeloid CC or lymphoid cell disorders, platelet disorders such as thrombocytopenia CC cand brain the and in tissue years and received as thrombocytopenia CC cand brain the and in tissue years and respective as the control and the research of bone, cartilage, tendon, ligament and/or nerve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAY-2000;
17-JUN-2000;
14-JUL-2000;
19-SEP-2000;
20-OCT-2000;
                                                                                                                                 and for regeneration of bone, cartilage, tendon, ligament and/or nerve tissue growth, and in tissue repair, healing of burns, incisions, ulcers, for treating osteoporosis, osteoarthritis, bone degenerative disorders, or periodontal disease. Furthermore, (I) is also useful for gut protection or regeneration and treatment of lung or liver fibrosis, reperfusion injury in various tissues, various immune deficiencies and disorders including severe combined immunodeficiency (SCID), bacterial or fungal infections, autoimmune disorders e.g. multiple sclerosis, reactions and conditions, such as asthma or other respiratory problems. In addition, (I) affects biorbythms or circadian cycles of rhythms, fertility, metabolism, catabolism, anabolism, storage or elimination of dietary fat, lipid, protein, carbohydrate, vitemins, minerals, provides analogs or effects or other respiratory problems.
analgesic effects or other pain reducing effects, immunoglobulin like activity and can act as an antigen in a vaccine composition to raise an immune response. AAS44920-AAS45295 represent novel human secreted proteicoding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel polypeptides and nucleic acids obtained from prepared from various human tissues, for diagnosis cancer, neurological, inflammatory, and autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   severe combined immunodeficiency; SCID; autoimmune multiple sclerosis; rheumatoid arthritis; diabetes fertility; analgesic; pain; antigen; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID
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; 2000US-0596193.
; 2000US-0616847.
; 2000US-0665363.
; 2000US-0693267.
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Drmanac
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           utoimmune disorder; allergy;
diabetes mellitus; asthma;
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Query Match Best Local S Matches 507

Similarity

94.9%; Score 507; I 100.0%; Pred. No. 1. tive 0; Mismatches

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Indels

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DB 22; Length 655;

Sequence

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RESULT 3
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ID AAH985
XX AAH985
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                     WPI; 200
P-PSDB;
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                                                                                                                                                                                                             25-JAN-2000; 2000US-0491404.
17-JUL-2000; 2000US-0617746.
03-AUG-2000; 2000US-0631451.
15-SEP-2000; 2000US-0663870.
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DB; AAM23856.
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A, Zhang
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Werhman T;
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Isolated polypeptide for treatment antibodies and research use -
                                                                of.
                                                                diseases,
                                                                diagnostics,
                                                                raising
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Claim 1275pp; English

The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs) from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a cDNA of the present sequence is a cDNA of the present sequence of the present sequence is a cDNA of the present sequence is a cDNA of the present sequence. the invention.

Sequence 538 BP; 141 A; 142 C; 115 G; 140 T; 0 other;

22;

Length

Similarity

á Ş 밁 Ś 밁 5 밁 S 밁 밁 S 밁 á В Ş 멍 δ 밁 Query Match Best Local S Matches 500 495 435 143 375 203 315 263 255 323 195 383 443 503 15 500; 23 83 75 CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAAACCACAACATTGACAG TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG GTGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGGGGCTCCGCCGCGGTGGCGGTTGCT TCTGTATTCTTCCAAAAAAAAAA 517 AAAAAGAAGTTTTGTAATTTATATTACTTTTTAGTTTTGATACTAAGTATTTAAACATATT TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT TCTGTATTCTTAAAAAAAAAAAA AAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATT TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT Conservative 93.3%; 0; Score 498.2; DB 2 Pred. No. 1.5e-99; Mismatches _ ω --Indels 0 Gaps 24 494 84 434 144 374 204 314 264 254 324 194 384 134 444

AAH98548 standard; cDNA; 538

12-OCT-2001 (first entry)

EST-derived coding sequence SEQ IJ S.

ARBSULT 4
AAH9845/C
ID AAH9855
XX AAH9855
XX AAH985
XX DT 12-OCT
XX Uman :
XX Human :
XX Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; tomato; monkey; dog; sea urchin; expressed sequence tag; EST diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition; ss.

Homo sapiens

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Query Match
Best Local S
Matches 500
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17-JUL-2000;
03-AUG-2000;
15-SEP-2000;
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Drmanac RA,
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¿A, Zhang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, cd diagnosis and treatment of diseases associated with inappropriate P cx expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions ci na patient's genome that affect the activity of P by expressing cc inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps. Cc Additionally, N may be used to produce the colon cancer-associated Ps. Dy inserting the nucleic acids into a host cell and culturing the cell co express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB7789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 499; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                present invention.

N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
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03-NOV-1999;
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This sequence represents human cDNA encoding

a secreted protein.

The

RESULT 7
AAF64012
ID AAF6
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AC AAF6
AC AAF6
AC AAF6
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DT 05-A
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AAF64012 standard;

cDNA;

ВP

05-APR-2001

(first entry) human secreted

cDNA encoding

Secreted protein; prevention; treatment; diagnosis; disease;

protein

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                identifying features of the cDNA sequences using a computer programme. The cDNAs are useful for expressing secreted proteins or fragments to obtain antibodies capable of specifically binding to the secreted proteins. The cDNAs may also be useful in diagnostic, forensic, gene therapy and chromosome mapping procedures and may be used to design expression vectors and secretion vectors. The proteins of the invention may be used to treat diseases including cancer, autoimmune diseases, cardiovascular disorders, cystic fibrosis, hypothyroidism, immunological disorders, obesity, mitocohondriocytopathies, diabetes, atherosclerosis, neurodegenerative disorders, graft rejection, Alzheimer's disease, dementia, hyperlipidaemia, septic shock and impotence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  secreted proteins. Also contained in the invention are methods for storing the sequence data on a computer system, and a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention relates to sequences AAA87725-A87774 which encode human secreted proteins AAB25763-B25812. The proteins include signal peption Included in the invention are a host cell containing one of the cDNA sequences, and a purified antibody capable of binding to one of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                     TCTGTATTCTTCCAAAAAAAAAAAAAAAA 522
                                                                                             AAAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTTTGATACTAAGTATTAAACATATT
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                                                                                                                                                                    TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAAGCCTGTGCATGAAA
                                                                     AAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                                                                                               TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
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RESULT
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Matches 505; (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 515 BP; 143 A; 106 C; 135 G; 130 T; 1 other;
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                                                                                                                                               TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
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                                                                                    AAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTTGATACTAAGTATTAAACATATT
                                                                                                                                 TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
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97US-0047583
97US-0047593
97US-0047593
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В δ 뮍 δ В S 밁 Ş 밁 S 망 Ś g Ś 용 S

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AAV59598 standard; DNA; 504
protein gene 88 clone HAUAV32
                                   entry)
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Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obseity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.

97US-0040336 97US-0040626 97US-0043311 97US-0043312 97US-0043313 97US-0043313 97US-0043568 97US-0043576 97US-0043576 97US-0040163 97US-0040161 98WO-US04493

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23-MAY-1997

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06-UIN-1997

06-UIN-1997

13-UIN-1997

13-UIN-1997

14-AUG-1997

22-AUG-1997

23-AUG-1997

23-AUG-1997

23-AUG-1997

24-AUG-1997

25-AUG-1997

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Kyaw
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This sequence represents a nucleic acid molecule designated Gene 88 the human cDNA clone HAUAV32 (deposited as clone ATCC 97901 and ATCC 209047) which encodes a secreted human protein. The gene can be use generate fusion proteins by linking to the gene to a human
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7 H,
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seful for
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                                                                                                                    immune
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e.g. cancers, neurological
  or blood disorders
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A, Greene JM, Hu JS
Olsen HS, Rosen C
Zeng Z;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunoglobulin Fc portion (e.g. AAV59502) for increasing the stability the fused protein as compared to the human protein only. The invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAW74731-W75026) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions ample or by determining the amount of the new polypeptides in a sample or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the 186 polynucleotides, based on which tissues they are most highly expressed (see AAV59511 for described uses).
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ABS73585
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                                                                                                                                                                                                                                                                                                                                                                                                                                CTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTG
standard;
                                                                                   TTCTTCCAAAAAAAAAAAAAAAAAA
                                                                                                                                          AAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTA
                                                                                                                                                                                                                                                                                             TCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
                                                                                                                                                                                                                                                                                                               TCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
                                                                                                                                                                                                                                                                                                                                                                      TGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGGGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCGAGCTGGGCGAGAAGTAGGGGAGGGC-ACGAGCCGCCGCGGTGGCGGTTGCTATCGCT
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                                                                   TTCTTCCAAAAAAAAAAAAAAAAAA
                                                                                                                         AAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTA
                                                                                                                                                                                GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
                                                                                                                                                                                                                                                                                                                                                     TGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTAT
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CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91.1%;
99.0%;
504
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Pred.
ВP
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No. 5.1
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RESULT 9
AB873585
IID AB873
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AC AB87
XX
DT 15-J
XX
DT 15-J
XX
DT Huma
XX
VA
KW Huma
                                                                         Human
                                                                                        15-JAN-2003
                                                                         CDNA
                                                                         #
                                                                                        (first
                                                                         for
                                                                         novel secreted
                                                                                        entry)
                                                                        protein gene
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cerebrovascular disorder; cerebral ischaemia; angiogenesis; sunburn; nervous system disorders; Alzheimer; s disease; infection; ocular disorder; corneal infection; wound healing; tissue regeneration; epithelial cell proliferation; organ transplantation; food additive; Human; ss; gene; secreted protein; autoimmune disease; chemotaxis; rheumatoid arthritis; hyperproliferative disorder; breast neoplasm; liver neoplasm cardiovascular disorder; cardiac arrest; skin aging;

88

ATCC used from

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08-SEP-1998;

16-JUL-2002.

US6420526-B1.

Homo

sapiens

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98US-0149476
        S-040333P

S-040333P

S-043312P

S-043312P

S-043312P

S-043312P

S-043569P

S-043569P

S-043569P

S-04750P

S-04750P

S-04750P

S-04759P

S-047612P

S-047612P
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3-040163P.
          The invention relates to an isolated protein that is one of 186 human CC secreted proteins, given in the specification, encoded by one of CC 309 cDNA sequences also given in the specification. The protein is used CC in a pharmaceutical composition used to prevent, treat or ameliorate a CC medical condition in e.g. humans, mice, rabbits, goats, horses, cats, CC dogs, chickens or sheep. Disorders which are diagnosed or treated include CC disorders e.g. neeplasms of the breast or liver, cardiovascular disorders CC e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, CC e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, CC e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, CC e.g. corneal infection. The polypeptides can also be used to aid wound CC healing and epithelial cell proliferation, to prevent skin aging due to subturn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to correct or decrease storage capabilities, fat content, lipid, protein, CC carbohydrate, vitamins, minerals, cofactors and other nutritional CC components. The present sequence represents a cDNA derived from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-AUG-1997
23-AUG-1997
23-AUG-1997
23-AUG-1997
23-AUG-1997
24-AUG-1997
25-SEP-1997
05-SEP-1997
05-SEP-1997
06-MAR-1998
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Duan R, Hu,
Moore PA, Sl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated human secreted protein for diagnosing, preventing, treating or ameliorating medical conditions and used as a food additive
                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-634796/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ruben
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         preservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lu J, Flo
Shi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rosen CA, Fischer CL, Soppet DP, Carter KC, Bec, Yu G, Ni J, Peng P, Young PE, Greene JM, Fel
hu J, Florence KA, Olsen HS, Ebner R, Brewer LA;
Shi Y, Lafleur DW, Li Y, Zeng Z, Kyaw H;
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENOME
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97US-056889P.
97US-056899P.
97US-056899P.
97US-056894P.
97US-056994P.
97US-056909P.
97US-056910P.
97US-056910P.
97US-056910P.
97US-056910P.
97US-056910P.
97US-056910P.
97US-05761P.
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97US-05761P.
97US-05764P.
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9718-05633P
9718-056662P
9718-056662P
9718-056662P
9718-056845P
9718-056862P
9718-056872P
9718-056874P
9718-056877P
9718-056876P
9718-056879P
9718-056882P
9718-056882P
9718-056882P
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Bednarik DR; Ferrie AM;

07-MAR-1997
07-MAR-1997
07-MAR-1997
07-MAR-1997
07-MAR-1997
07-MAR-1997
11-APR-1997
12-MAY-1997
13-MAY-1997
13-JUN-1997
13-JUN-1997
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13-JUN-1997
13-JUN-1997
13-JUG-1997
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13-JUG-1997
13-JUG-1997

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RESULT 10
AAV59746
ID AAV59746
AC AAV59
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Best Local S
Matches 500
                                                                                                                                                                                        Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
                                                                                                                                 cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene encoding one of the novel human secreted proteins of the invention. Note: This sequence did not form part of the printed specification, but was obtained in electronic format directly from USPTO at sequence.html?DocID=6420526B1.
                                                                                                                                                                                                                                                                                                                                                                                        AAV59746;
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                    11-SEP-1998
                                                          WO9839448-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
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                                                                                                                                                                                                                                                                                                             protein gene
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99.0%;
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Pred. No. 5.1
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                                                                                                                                                                          which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polymocleotides. Specific uses are described for each of the 186 polynucleotides, based on which tissues they are most highly expressed in (see AAV59511 for described uses).
                                                                                                                                                                                                                                                                                    This sequence represents a nucleic acid molecule designated Gene 88 from the human cDNA clone HAUAV32 (deposited as clone ATCC 97897 and ATCC 209043) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin Fc portion (e.g. AAVS9502) for increasing the stability of the fused protein as compared to the human protein only.

The invention relates to 186 novel genes and their fragments (nucleic the invention relates to 186 novel genes and their fragments)
                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
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Kyaw
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GGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGGTCTGCAGACGCGATGGA
                                          GCAGGATTTGGCCAGGACCCSCCGCGGTGGCGGTTGCTATCGCTTCGCAGAACCTACTCA
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                                                                                                                                                                                                                                                                                                                                                              angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, minerals, cofactors and other nutritional components. The present sequence represents a cDNA derived from a gene encoding one of the novel human secreted proteins of the invention. Note: This sequence did not form part of the princed specification, but was obtained in electronic format directly from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated protein that is one of 186 human secreted proteins, given in the specification, encoded by one of 309 cDNA sequences also given in the specification. The protein is used in a pharmaceutical composition used to prevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated human secreted protein for diagnosing, preventing, treating or ameliorating medical conditions and used as a food
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P-PSDB; ABG95421.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or preservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Endress GA, Y
Duan R, Hu J,
                                                                                                                                                                                                                                                                                                               Sequence 506 BP; 148 A; 103 C; 121 G; 132 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID No
                                                                                                                                                                                                                                                                                                                                                 seqdata.uspto.gov/sequence.html?DocID=6420526B1.
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                                                                  157
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GATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATC
                                                                                                                         GCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCTGCTGCAGACGCGATGGA
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                                                       TAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAA
                                                                                                      GGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCGGTCTGCAGACGCGATGGA
                                                                                                                                                                         GCAGGATTTGGCCAGGACCCSCCGCGGTGGCGGTTGCTATCGCTTCGCAGAACCTACTCA
                                  TAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAA
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A, Yu G, Ni J, Feng P, Young PE,
Hu J, Florence KA, Olsen HS, Ebner
Shi Y, Lafleur DW, Li Y, Zeng Z,
                                                                                                                                                                                                                                              Conservative
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97US-057611P.
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97US-057650P.
97US-057650P.
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97US-056903P.
97US-056908P.
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                                                                                                                                                                                                                                          Score 478.6; 1
Pred. No. 2.8e
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Young PE, Greene JM,
n HS, Ebner R, Brewer
Y, Zeng Z, Kyaw H;
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nes 11;
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RESULT 12

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13-NOV-1997;
17-DEC-1997;
                                      AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used to express secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures. Uses also include design of expression vectors and secretion vectors.
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  Sequence
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13-APR-1998;
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RESULT 13
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                                                                                                                                                                                                                                                                                                          multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis allergic reaction; osteoporosis; osteoarthritis; periodontal disease; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; liver fibrosis; lung fibrosis; reperfusion injus systemic cytokine damage; tissue differentiation; contraceptive; strol coagulation disorder; myocardial infarction; inflammatory condition; septic shock; sepsis; ischaemia; reperfusion injury; arthritis; tumous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human protein; hydrophobic domain; nutritional source; haematopoiesis; cytokine production; cell proliferation; cell differentiation; immune deficiency; infectious disease; autoimmune disorder; asthma;
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Pred. No. 1.1e
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24-JUL-1998; 07-AUG-1998;

98JP-0208820 98JP-0224105 99WO-JP03929

22-JUL-1999;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other cytokines in concern from the control of various immune stimulating or immune suppressing activity. It can be used intreatment of various immune deficiencies and disorders, and to treat treatment of various immune deficiencies and disorders, and to treat treatment of various answers of the control o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence encodes a human protein of the invention, which has hydrophobic domains. The DNA sequences can be used as a probe or as a genetic marker. The protein can also be used as a marker, and to identify potential genetic disorders. The DNA and protein can also be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other cytokines in certain cell populations. The protein also exhibits
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09-SEP-1998;
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TGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTG
                                                                                                                                                                                                                                                                                                                                              AGGGGAGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTCGCAGAACCTACTCAGG
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                                                                                               ACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAAGA
                                                                                                                                                                                                                     CAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCGGTCTGCAGACGCGATGGATA
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98JP-0254736.
98JP-0275505.
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                                                     TGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATC
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                                                                                                                                                                                       TGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTG
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                                                                              458
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ABK36006 standard; cDNA; 439

cDNA sequence #397 encoding novel human secreted protein

08-MAY-2002

(first

entry)

immune deficiency disorder; blood disorder; inflammatory disorder; infectious disorder; allergic condition; neurodegenerative disorder; liver fibrosis; coagulation disorder; gene therapy; antimicrobial; Human secreted cancer; hepatotropic; protein; hyperproliferative disorder; autoimmune immunosuppressive; antirheumatic; gene; ss disorder;

Homo sapiens

18-OCT-2001

29-MAR-2001; 2001WO-US10232

06-APR-2000; 2000US-195605P

_ GENETICS INST INC.

Jacobs K, M Merberg D, Clark HF, McCoy JM, Treacy M Treacy M, Fechtel K, Lavallie Agostino N Howes SH, o MJ, bu. Resnick F ER Collins-Racie LA, Evans C; Bowman MR, Spaulding V, RJ, Gulukota Graham Wong GG; caham JR;

WPI; 2002-179322/23.

Six hundred and twenty three polynucleotides derived from a variety of human tissue sources which encode secreted proteins, useful for treating immune deficiencies and disorders such as autoimmune disorder disorders

Claim ۳. Page 296-297; 393pp; English

RESULT 14
ABK36006
ID ABK36
XX ABK36
XX ABK36
XX DE CDNA
XX Human
XW Iinfec
XW Iinfec
XW Iinfec
XW Itwo
XX Homo
XX Hom sclerosis), blood disorders (e.g. LILLINGO), infectious disorders (e.g. hepatitis), disorders (e.g. arthritis), infectious disorders (e.g. disorders (e.g. allergic conditions (e.g. asthma), neurodegenerative disorders (e.g. alrheimer's disease), liver fibrosis, coagulation disorders (e.g. which encode human secreted proteins. The cDNA sequences have been derived from a variety of human tissues. The invention also provides a method for producing proteins from these polynucleotide sequences. The proteins are useful for identifying compounds that modulate their activity and production. The sequences of the invention are useful for treating diseases such as hyperproliferative disorders (e.g. cancer), immune deficiency disorders (e.g. severe combined immunodeficiency (SCID)), autoimmune disorders (e.g. multiple sclerosis), blood disorders (e.g. thrombocytopaenia), inflammatory invention relates to the isolation of novel cDNA sequences

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AAX997373
ID AAX977
XX AAX97
AC AAX97
XX AAX97
XX BECTE
XX SECTE
XX GAIAGI
COS HOMO
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17-DEC-1997;
09-FEB-1998;
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10-AUG-1998;
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                        98US-0099273.
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Pred. No. 1.4e-83;
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Best Local Similarity
Matches 412; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used texpress secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures. Uses also include design of expression vectors and secretion vectors.
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Pred. No. 1.3e-80;
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Search completed: January time 31, 2004, 02:59

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

4	w	N	س	Result No.
501.8	505.8	507.4	510.2	Score
94.0	94.7	95.0	95.5	Query Match
519	524	552	548	Query Match Length DB
13	10	14	13	DB
BX091668	BG705303	CB993661	BU532309	Query Match Length DB ID
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W52820 W38899 AA305052	AI141284	AA989129	AA911088	AI989739	AW183193	AA516431	AI989747	BM553628	AI826623	BF215121	BU536995	BG532587	AV759888	BG613984	AI128804	AA455042	BU564679	AA429945	AV754613	BU566814	AI743235	BX436125	AI078580	BU533237	BF203359 ·	BF109912	BF691818	BU189908	BU565721	BX464689	BX422104	BG031757	6649	7208	G5290	œ	G24999	U5992
W38899 ZC35CUB. F1 W38899 ZB32G04.F1 AA305052 EST176055	34 qa52h10.	29 or75h10.	AA911088 ok67e01.s	wu21e1	AW183193 xj67f11.x	AA516431 ne58a03.s	AI989747 wu21f09.x	BM553628 AGENCOURT	AI826623 wk35d10.x	BF215121 601846829	BU536995 AGENCOURT	BG532587 602562152	AV759888 AV759888	BG613984 602641513	AI128804 qa94b08.s	AA455042 aa04a07.s	BU564679 AGENCOURT	AA429945 zw67f10.s	AV754613 AV754613	BU566814 AGENCOURT	AI743235 wg90a02.x	BX436125 BX436125	_	BU533237 AGENCOURT	Ψ		w	w	-	w	_	BG031757 602299819	_	٠,	9072	F399486	9994 60236206	U599247 AGENCOU

ALIGNMENTS

	COMMENT	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BU532309	D 1113 13
Email: cgapbs r@mail.nih.gov Tissue Procurement: NCI CDNA Library Preparation: Michael Brownstein Laboratory cDNA Library Preparation: Michael Brownstein Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM2726 row: i column: 16 High quality sequence stop: 516.	Contact: Robert Strausberg, Ph.D.	National Institutes of Health, Mammalian Gene Collection (MGC)	NIH-MGC http://mgc.nci.nih.gov/.	1 (bases 1 to 548)	Eukarvota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	Homo sapiens (human)	EST.	BU532309.1 GI:22842750	BU532309	IMAGE:6557848 5', mRNA sequence.	AGENCOURT_10202846 NIH_MGC_126 Homo sapiens cDNA clone	BU532309 548 bp mRNA linear EST 13-SEP-2002		

information can

be

(NHGRI) with help

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                                                                                                                                                                                                                                                                                                                                                                                                                          TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
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                                                                                 TCTGTATTCTTCCAAAAAAAAAAAAAAAAAAATTANNNAA
                                                                                                             AAAAAGAAGTTTTGTAATTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATT
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                                                                                                                                                                                                                                            TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA 434
                                                                                                                                                                                                                                                                            TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGCCCCTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)NN-3'. Full-length enriched library was constructed using the Clontech Creator SMART kit and size-selected to contain the 0.5-1 kb size fraction (other fractions present in NIH_MGC 127 and NIH_MGC 128). Library created in the laboratory of T. Usdin, M.D., Ph.D. (NIMH, NIH). Note: this is a NIH_MGC Library."
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Pred. No. 3.3e-49;
0; Mismatches 8
   552
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   mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Stefan Hansson
cDNA Library Preparation: Michael J. Brownstein (NHGRI) w
and advice from Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: NDAM353 row: a column: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 552)
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                                                                                                                                                                                                 CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCCTTCTGCT
                                                                                                                                                                                                                                                                  ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAAGAGTTGAGGGAAAGTGCTGCTG
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TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                  TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGGTGGATATTATCAACTCACTGGTAACAA
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/tissue_type="pre-eclamptic placenta"
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/db_xref="taxon:9606"
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Pred. No. 6.8e-49;
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI),
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BG705303 524 UP 602687808F1 NIH_MGC_95 Homo sapiens
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High quality sequence stop: 521.
Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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BG705303.1 GI:13979504
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
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Clone distribution: MGC clone distribution information can
cound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
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  ATCGCTTCGCAGAACCTACTCAGGCAGCCGAGCTGAGAAGAGTTGAGGGGAAAGTGCTGCTG 134
                                         GGGAGGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                     GTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
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/Clone_lib="NIH_MGC_95"
/note="Organ: brain; Vector: pBluescriptR (modified / note="Organ: brain; Vector: pBluescriptR (modified / note="Organ: brain; Vector: pBluescriptR (psine) |
pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size_2: Skb and normallized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

13 a 113 c 138 g 131 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:4820568"
/tissue_type="hippocampus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
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                                                                                                                                             94.7%;
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Pred. No. 1.1e-48;
0; Mismatches 2;
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                                                                                                                                www.rzpd.de
This clone is available royalty-free from RZPD;
This clone (Connewrzpd.de) for further information.
Contact RZPD (Clone@rzpd.de) for further information.
M13r, Primer sequence: TTTCACACAGGAAACAGCTATGAC.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 519)

Ebert, L., Heil, O., Hennig, S., Neubert, P., Partsch, E., Peters, M., Radelof, U., Schneider, D. and Korn, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BX091668 Soares ovary tumor NbHOT Homo sapiens IMAGE998M041818 ; IMAGE:740523, mRNA sequence.
                                                                                                                                                                                                                                                Heubnerweg 6, D-14059 Berlin,
Tel: +49 30 32639 101
Fax: +49 30 32639 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
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                                                                                                                                                                                                                                                                                             bin/showLib.pl.cgi/response?libNo=972 Contact: Ina Rolfs
RZPD Deutsches Ressourcenzentrum fuer Genomforschung Gmbl
Heubnerweg 6, D-14059 Berlin, Germany
                                                                                                                                                                                                                                                                                                                                                                              RZPD; IMAGp998M041818.
RZPDLIB; I.M.A.G.E. cDNA Clone Collection;
Human UnigeneSet - RZPD3 (RZPDLIB No.972)
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Im Neuenheimer Feld 580, D-69120 Heidelberg, Germany
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                                                             /mol_type="mRNA"
/db_xref="taxon:9606"
                   /clone="IMAGp998M041818 ; IMAGE:740523"
/sex="Female"
tissue_type="ovarian tumor"
                                                                                                    organism="Homo sapiens"
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EST 23-JAN-2003 clone

363

374 303 314 243 254 183

483 494 423 434

GmbH

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JOURNAL COMMENT
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                                                                                                                                                        AGENCOURT 8908732 NIH MGC 142 Homo sapiens 5', mRNA sequence.
Unpublished
Contact: Robert Strausberg, Ph.
Email: cgapbs-r@mail.nih.gov
                                       1 (bases 1 to 528)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, N
                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                     Homo sapiens (human)
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Location/Qualifiers
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CDNA Library Preparation: Michael Brownstein Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGAAGCCGAGCTGGGGGAGAAGTAGGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
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                                                          AAAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTTGATACTAAGTATTAAACATATT
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5'-AAGCAGTGCTATCACAGCAGAGTGGCCAGTATACGGCCGGG-3' and
5'-ATTCTAGAGGCCGAGAGTGGCCGACATG-dT(30)NN-3'. Full-length
enriched library was constructed using the Clontech
Creator SMART kit and size-selected to contain the >0.5 kb
size fraction (other fractions present in NHI_MGC_141).
Library created in the laboratory of M. Brownstein (NIMH,
NIH). Note: this is a NIH_MGC Library."

107 c 134 g 131 t 26 others
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/note="Vector: pDNR-LIB; Site 1: SfiI (ggccattatggcc);
/note="Vector: pDNR-LIB; Site 1: SfiI (ggccattatggcc);
SfiI (ggccgctcggcc); Double-stranded cDNA was prepared from a pool of 40 cell line polyA+ RNAs (bladder 2%, blood - 33.4%, brain - 5.6%, breast - 12.5%, colon - 2%, blood - 33.4%, brain - 5.6%, breast - 12.5%, colon - 4%, connective tissue - 1.4%, eye - 1%, intestine - 2.6%, kidney - 2.2%, liver - 5.7%, lung - 10.8%, NK-cell - 5.2%, ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, prostate - 4.3%, salivary ovary - 4%, pharyux - 2.5%, ph
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/lab host="DH10B (T1-phage-resistant)"
/clone_lib="NIH_MGC_142"
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/db_xref="taxon:9606"
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Pred. No. 3.5e-47;
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1 (Dases 1 to 962)

NIH-MGC http://mgc.nci.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
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                                                                                          CACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTC
                                                                                                                                                                                GCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGC
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                                ATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTG
                                                                                                                                              GCGATGGATAACGTGCAGCCGAAAATAAAAACATCGCCCCTTCTGCTTCAGTGTGAAAAGGC
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                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="adenocarcinoma, cell line"
/lab_host="DH10B (phage resistant)"
/clone_lib="NIH MGC_90"
/clone_Tib="NIH MGC_90"
/note="Organ: liver; Vector: pCMV-SPORT6; Site 1: Not1;
/note="Organ: liver; Vector: pCMV-SPORT6; Site 1: Not1;
/note="Sall; Cloned unidirectionally; oligo-dT primed.
Average insert size 1.7 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
a 199 c 288 g 141 t
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                clone="IMAGE:4470662"
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Pred. No. 2.3e-47;
0; Mismatches 1;
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide that was used to prime the synthesis of first
strand cDNA and therefore this may represent a bonafide poly A
tail. cDNA Library Preparation: M.B. Soares Lab Clone distribution:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                שיטיים אינים איני
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EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: bento-soares@uiowa.edu
The sequence contained an oligo-dT track that was present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 319 335 8250 Fax: 319 335 9565
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375 Newton Road , 4156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact:
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                             130
                                                                 Soares, MB
                             TAG_SEQ=None
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/clone="UI-R-CA1-bjb-b-12-0-UI"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  strain="Sprague-Dawley"
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Best Local Similarity 99.6
Matches 493; Conservative
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                                                              cDNA Library Preparation: CLONETECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. CONSORTIUM (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLCM1556 row: i column: 18
                                                                                                                                                      Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                              BG529072 /bI
602579169F1 NIH_MGC_60 Homo
                                                                                                                                                                                                                                                                                                                          mRNA sequence.
BG529072
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National Institutes of Health,
                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                 BG529072.1 GI:13520609
                                                                                                                                                                                             Unpublished
                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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                                                                                                                                                                                                                                   (bases 1 to 781)
                                                   quality sequence stop: 475.
/organism="Homo sapiens"
/mol_type="mRNA"
                                     ocation/Qualifiers
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Pred. No. 4.2e-47;
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mo sapiens
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                           Homo sapiens (human)
                                                      BM472086.1
EST.
                                                                                         AGENCOURT_6469731 NIH_MGC_92 Homo 5', mRNA sequence.
                                                                                                                     BM472086
   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/tissue_type="denocarcinoma"
/lab_host="DH10B (TI phage-resistant)"
/lab host="DH10B (TI phage-resistant)"
/clone_lib="NIH_MGC_60"
/clone_lib="NIH_MGC_60"
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
/note="Organ: prostate from cell line RNA.
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Pred. No. 1.1e-46;
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM12328 row: a column: 12
High quality sequence stop: 486.
Location/Qualifiers
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Tissue Procurement: ATCC
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National Institutes of Health, Mammalian
                                                                                          AGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTAT 275
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/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone="IMAGB:5576339"
/tissue_type="embryonal carcinoma, cell line"
/tissue_type="embryonal carcinoma, cell line"
/tlstue_type="embryonal carcinoma, cell line"
/lab_host="NHIDB (phage-resistant)"
/clone_lib="NHIMGC 92"
/clone_site="Organ: testis; Vector: pCMV-SPORT6; Site_1: NotI;
/note="Organ: testis; Vector: pCMV-SPORT6; Site_1: NotI;
/site_2: SalI; Cloned unidirectionally; oligo-dT primed.
Average insert size 2.5 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
135 a 103 C 121 g 128 t
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National Institutes of Health,
Unpublished
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: NCI
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 526)
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AGENCOURT 10397961 NIH MGC 141 Homo
IMAGE:6605733 5', mRNA sequence.
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                                                                                                                                                                                          GCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCTGGTTCT
                                                                                                                                                                                                                                                                                    GAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTC
AAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTC
                                                                                                    GCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTG
                                                                                                                                                                  GCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGGTCT
                                                                                                                                                                                                                                                          GAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTC
                                                                         GCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTG
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5'-AACCAGTGGTATCAACGCAAGTGGCCATTACGGCCGG-3' and 5'-AATTCTAGAGGCCGACGAGAGTGGCCAATTACGGCCGG-3' and 5'-AATTCTAGAGGCCGACGAGAGTGGCCCATTACGGCCGG-3' and 5'-AATTCTAGAGGCCGACGAGTGGCCCATTACGGCCGGG-3' and 5'-AATTCTAGAGGCCGACGGCGAATG-dT(30)NN-3'. Full-length enriched library was constructed using the Clontech Creator SMART kit and size-selected to contain the 0.2-0.5 kb size fraction (other fractions present in NIH MGC 142). Library created in the laboratory of M. Brownstein (NIMH, NIH). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="mixed (pool of 40 RNAs)"
/lab_host="DH10B (T1-phage-resistant)"
/clone_lib="WIH_MGC_141"
/note="Vector: \bar{p}DNR-LIB; Site_1: SfiI (ggccattatggcc);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/db_xref="taxon:9606"
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95.9%;
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Pred. No. 2.2e-46;
0; Mismatches 21
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                                                   499;
 30 GCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTCGCAGAAC 89
                                                                                                                                                                                                                                                                                                                                                                        found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLAM10089 row: e column: 06 High quality sequence stop: 446.
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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BG031757
BG031757.1 GI:12422364
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602299819F1 NIH_MGC_87 Homo sapiens cDNA clone IMAGE:4394093 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                Similarity
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                                                   Conservative
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                                                                                                                              /tissue_type="mammary adenocarcinoma, cell line"
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/clone_top="NIH_MGC_87"
/note="Togan: breast; Vector: pCMV-SPORT6; Site_1: NotI;
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/site_2: SalI; Cloned unidirectionally; oligo-dT_primed.
Average insert size 1.383 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC_Library."
a 108 c 132 g 131 t
                                                                                                                                                                                                                                                                         /mol_type="mRNA"
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/clone="IMAGE:4394093"
                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
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/clone="CSODHOO7YB16"
/tissue_type="T_CELLS_(JURKAT_CELL_LINE)"
/cell_line="JURKAT_CELL_LINE"
/cell_lib="Homo sapiens T_CELLS_(JURKAT_CELL_LI
/note="Vector: pCMVSPORT_6; lst strand_cDNA_was
                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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sequence

InVitroGen Corporation 1600 ID : CSODH007DA08QP1.

LINE) " primed

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Mammalia; Eutheria; Primate
1 (bases 1 to 671)
Li,W.B., Gruber,C., Jessee,
Full-length cDNA libraries
Unpublished
http://www.genoscope.cns.fr/
cgi-bin/cluster.cgi?seq=CSODHOO7DAO8QP1&cluster=8312.f. Contact
Feng Liang Email: fliang@lifetech.com URL:
http://fulllength.invitrogen.com/ InVitroGen Corporation 1600
                                                                                                         Email: segref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen. This sequence belongs to sequence cluster 8312.f For
more information about this cluster, see
                                                                                                                                                                                                                          Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
                                                                                                                                                                                                                                                                             Contact: Genoscope
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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           Contact: Genoscope
Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen. This sequence belongs to sequence cluster 8312.f
more information about this cluster, see
http://www.genoscope.cns.fr/
                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vert. Mammalia; Eutheria; Primates; Catarrhini; Hol (bases 1 to 1201)
Li, W.B., Gruber, C., Jessee, J. and Polayes, D. Full-length cDNA libraries and normalization Unpublished
                                                                                                                                                                                                                                                                                                             BX464689 Homo sapiens T CELLS (JURKAT CELL LINE) clone CSODHOO7YB16 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                     Homo
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cgi-bin/cluster.cgi?seq=CS1DH002ZF11QP1&cluster=8312.f. Contact
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Pred. No. 2.8e-45;
4; Mismatches 5
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SOURCE ORGANISM VERSION KEYWORDS

Homo sapiens Homo sapiens (human)

Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

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RESULT 14
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AGENCOURT 10401193 NIH MGC 141 Homo IMAGE: 6604530 5', mRNA sequence. BU565721.1 GI:22916011 EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Feng Liang Email : fliang@lifetech.com URL : http://fulllength.invitrogen.com/ InVitroGen Corporation 1600 Faraday Avenue Genoscope sequence ID : CS1DH002ZF11QP1.
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/note="Vector: pcMvSpORT 6; 1st strand cDNA was primed
with a NotI-oligo(dT) primer. Five prime end enriched,
double-strand cDNA was digested with Not I and cloned i
the Not I and ECORV sites of the pCMvSpORT 6 vector.
Library was not normalized."
173 others
a 298 c 248 g 180 t 173 others
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/tissue_type="T CELLS (JURKAT CELL LINE)
/cell_line="JURKAT CELL LINE"
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.ni
Tissue Procurement: NCI
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                                                                                                                                     GACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCC 399
                                                                                                                                                                                                                                                                                                                                              GCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGT
GTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGT
                                                                                                  GACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCC
                                                                                                                                                                                                                                                           GTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGGTGTTTGCACTTGT
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/clone type="mixed (pool of 40 RNAs)"
/lab_host="DH10B (T1-phage-resistant)"
/clone lib="NHH MCC 141"
/clone lib="NHH MCC 141"
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/clone lib="NHH MCC 142"
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Pred. No. 1.3e-44;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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AGENCOURT 8036927 NIH MGC 72
5', mRNA sequence.
BUI 80000
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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                                                                 ACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAAGA
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                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                            /tissue_type="melanotic melanoma"
/lab host="DH10B (phage-resistant)"
/clone_lib="NHH MGC 72"
/note="Organ: skin; Vector: pcMV-SPORT6; Si
Site_2: Sall; Cloned unidirectionally. Pri
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/db_xref="taxon:9606"
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Search completed. January 31 2004 04:09:42	421 TTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCACATA 474		361 CCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAA	399	301 TGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATC 360		241 TGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTG 300	279

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Result
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1: /cgn2_6/ptodata/2
2: /ggn2_6/ptodata/2
3: /cgn2_6/ptodata/2
4: /cgn2_6/ptodata/2
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
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US-09-663-600A-53
US-09-620-132-147
US-09-633-600A-147
US-09-599-360B-70
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US-08-956-976-2
US-09-256-976-2
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Compugen Ltd
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Sequence 98, Appl
Sequence 252, Appl
Sequence 53, Appl
Sequence 164, App
Sequence 147, App
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95, Appl
12, Appl
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64, Appl
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8.1 3238 3 928 8.1 3238 2 928 8.0 1332 2 93 94 94 97 97 97 97 97 97 97 97 97 97 97 97 97
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US-US-123-934A-5 US-09-123-934A-5 US-09-057-762-1 US-09-057-762-1 US-09-057-762-7 US-09-795-927-5 US-09-795-927-5 US-09-7149-476-316 US-09-149-476-193 US-09-149-475C-196 US-09-328-475C-74

ALIGNMENTS

RESULT 1 US-09-599-360B-29

Sequence 29, Application US/09599360B Patent No. 6548633
GENERAL INFORMATION:

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SEQ ID NO 29
                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                         Matches
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APPLICANT: Bougueleret, L.
APPLICANT: Jobert, S.
TITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides
FILE REFERENCE: GENSET.050CP3
CURRENT APPLICATION NUMBER: U5/09/599,360B
CURRENT FILING DATE: 2000-06-21
PRIOR APPLICATION NUMBER: 60/113,686
PRIOR FILING DATE: 1998-12-22
PRIOR APPLICATION NUMBER: 60/141,032
PRIOR APPLICATION NUMBER: 60/141,032
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 123
                                                                                                                                                                                                                                                           NAME/KEY: polyA_signal
LOCATION: 457..462
NAME/KEY: polyA_site
LOCATION: 500..515
NAME/KEY: misc_feature
                                                                                                                                                                                                                            OTHER INFORMATION: n=a,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo
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LOCATION: 144..287
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LOCATION: 144..440
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ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAAGAGTTGAGGGAAAAGTGCTGCTG 134
                                                                GTGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGTGCTCCGCCGCCGGTGGCGGTTGCT
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Pred. No. 1.1e-116;
0; Mismatches 2;
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RESULT 2 US-09-149-476-98 Sequence 99, Application US/09149476. Patent No. 6420526 Patent No. 6420526 Patent No. 6420526 Patent No. 6420526 Patent No. 16420526 Patent No. 16420526 Patent No. 16420526 Patent Pilla North	Db 67 ATCGCTTCGCAGAACCTACTCAGGCAGCAGAAAAAAAAAA
EARLIER APPLICATION NUMBER: 60/043,568 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,311 EARLIER APPLICATION NUMBER: 60/043,671 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,674 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,312 EARLIER FILLING DATE: 1997-04-11 EARLIER FILLING DATE: 1997-04-11 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,312 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,313 EARLIER FILLING DATE: 1997-04-11 EARLIER FILLING DATE: 1997-04-11 EARLIER APPLICATION NUMBER: 60/043,315 EARLIER APPLICATION NUMBER: 60/043,315 EARLIER APPLICATION NUMBER: 60/056,867 EARLIER FILLING DATE: 1997-08-22 EARLIER APPLICATION NUMBER: 60/056,897 EARLIER APPLICATION NUMBER: 60/056,897 EARLIER APPLICATION NUMBER: 60/056,897 EARLIER APPLICATION NUMBER: 60/056,897 EARLIER APPLICATION NUMBER: 60/056,630 EARLIER FILLING DATE: 1997-08-22 EARLIER PILLING DATE: 1997-08-22 EARLIER APPLICATION NUMBER: 60/056,630 EARLIER APPLICATION NUMBER: 60/056,637 EARLIER APPLICATION NUMBER: 60/056,637 EARLIER FILLING DATE: 1997-08-22 EARLIER FILLING DATE: 1997-08-22 EARLIER APPLICATION NUMBER: 60/056,637 EARLIER APPLICATION NUMBER: 60/056,637 EARLIER FILLING DATE: 1997-08-22 EARLIER FILLING DATE: 1997-08-22 EARLIER APPLICATION NUMBER: 60/056,662 EARLIER FILLING DATE: 1997-08-22 EARLIER APPLICATION NUMBER: 60/056,662 EARLIER FILLING DATE: 1997-08-22	RILING DATE: 1997-05-23 R APPLICATION NUMBER: 60/047, R FILING DATE: 1997-05-23

APPLICATION NUMBER: 60/056,637
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,903
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,888

1997-08-22

N NUMBER: 60/UE: 1997-08-22

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FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,874 FILING DATE: 1997-08-22

APPLICATION NUMBER: 60/056,636 APPLICATION NUMBER: 60/056,911 FILING DATE: 1997-08-22

APPLICATION NUMBER: 60/056,910

1997-08-22

APPLICATION NUMBER: 60/056,894 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,880 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,879

FILING DATE:

1997-08-22

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Sequence 252, Application US/09149476
PAtent No. 6420526
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: 186 Human Secreted protei
FILE REFERENCE: PZ002P1
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: PCT/US98/04493
EARLIER FILING DATE: 1998-03-06
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER APPLICATION NUMBER: 60/040,333
EARLIER APPLICATION NUMBER: 60/040,333
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ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/057,669
ER FILING DATE: 1997-09-05
ER APPLICATION NUMBER: 60/049,610
ER FILING DATE: 1997-06-13
ER APPLICATION NUMBER: 60/061,060
ER FILING DATE: 1997-10-02
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R APPLICATION NUMBER: 60/0:
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/0:
R FILING DATE: 1997-66-06
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Pred. No. 5.9
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EARLIER

APPLICATION NUMBER: 60/047,595 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,599 FILING DATE: 1997-05-23

APPLICATION NUMBER: 60/057,761

APPLICATION NUMBER: 60/056,892

1997-08-22 1997-08-22

60/056,845

FILING DATE:

1997-08-22

FILING DATE: 1997-08
APPLICATION NUMBER:

FILING DATE:

APPLICATION NUMBER: 60/056,631

FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,864 FILING DATE: 1997-08-22

FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,586 FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,585 APPLICATION NUMBER: 60/047,588 FILING DATE: 1997-05-23

APPLICATION NUMBER: 60/047,590

1997-05-23

FILING DATE:

APPLICATION NUMBER: 60/047,594

1997-05-23

LING DATE: 1997-05-23

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R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056,881
R FILING DATE: 1997-08-22
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R PILING DATE: 1997-08-22
R FILING DATE: 1997-08-22
R FILING DATE: 1997-08-22

APPLICATION NUMBER: 60/056,632 FILING DATE: 1997-08-22

APPLICATION NUMBER: 60/043,670 FILING DATE: 1997-04-11

FILING DATE: 1997-05-23

FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,876

APPLICATION NUMBER: 60/056,664

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APPLICATION NUMBER: 60/056,862 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,887

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FILING DATE:
APPLICATION |

FILING DATE:

NUMBER: 60/043,578: 1997-04-11

1997-05-23

APPLICATION NUMBER: 60/047,589 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,593

FILING DATE:

FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,614

APPLICATION NUMBER: 60/043,576
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/047,501

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ER APPLICATION NUMBER: 60/038,621
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,334
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,163
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,163
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/047,600
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,507
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,507
ER APPLICATION NUMBER: 60/043,507
ER APPLICATION NUMBER: 60/043,507
ER APPLICATION NUMBER: 60/043,607
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,607
ER FILING DATE: 199

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ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,876
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APPLICATION NUMBER: 60/056,884
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/057,669
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APPLICATION NUMBER: 60/061,060
FILING DATE: 1997-10-02
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APPLICATION NUMBER: 60/056,875
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,862
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APPLICATION NUMBER: 60/048,964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/056,908
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APPLICATION NUMBER: 60/056,887
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                                                                                                                                                                                                                                                                                                 GATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATC
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                                                                                                                                  TGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAA
                                              TGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACT
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Pred. No. 6.4e-113;
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PRIOR FILING DATE: 1997-11-13
PRIOR PELING DATE: 1997-12-17
PRIOR PELING DATE: 1997-12-17
PRIOR PELING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR PILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 229
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Patent No. 6573068
                                                                                                                                                              NAME/KEY: polyA signal
LOCATION: 464.469
NAME/KEY: polyA site
LOCATION: 489.500
NAME/KEY: misc_feature
LOCATION: 197.412
OTHER INFORMATION: id: AA429945
OTHER INFORMATION: est
NAME/KEY: misc_feature
LOCATION: 61.195
OTHER INFORMATION: homology
OTHER INFORMATION: homology
OTHER INFORMATION: di: AA429945
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SEQ ID NO 53
LENGTH: 500
                    NAME/KEY: misc_feature LOCATION: 197..412
OTHER INFORMATION: homol OTHER INFORMATION: id :A OTHER INFORMATION: est NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: Sig_peptide
LOCATION: 199..288
OTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 5.599999046
OTHER INFORMATION: seg IVSVLALIPETTT/
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NAME/KEY: misc_feature
LOCATION: 339..401
OTHER INFORMATION: homol
OTHER INFORMATION: id: P
OTHER INFORMATION: est
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LOCATION: 197.324
OTHER INFORMATION: homo:
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OTHER INFORMATION: est
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OTHER INFORMATION: 1
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OTHER INFORMATION: 6
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LOCATION: 197._338
OTHER INFORMATION: homo:
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LOCATION: 207..412
OTHER INFORMATION: homology
OTHER INFORMATION: id :W93646
OTHER INFORMATION: est
                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: 71.155
OTHER INFORMATION: homology
OTHER INFORMATION: id :W52820
OTHER INFORMATION: est
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LOCATION: 425._469
OTHER INFORMATION: homo
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LOCATION: 58..195
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CACAMION: 425..488
THER INFORMATION: homology
THER INFORMATION: id: AA516431
THER INFORMATION: est
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JOCATION: 52..195
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INFORMATION: id :AA516431
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id :AA455042
homology
id :W52820
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US-09-220-132-164/c
; Sequence 164, Application US/09220132
; Patent No. 6506607
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                                                   NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 164
LENGTH: 526
                                                                                                                                                                                                                                                                                             GENERAL INFORMATION: APPLICANT: Shyjan,
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Best Local 9
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OTHER INFORMATION: id: w19506
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 9..10,12
OTHER INFORMATION: n=a, g, c or
                                                                                                                      APPLICANT: Shyjan, Andrew W.

TITLE OF INVENTION: METHODS AND COMPOSITONS FOR THE IDENTIFICATION ITITLE OF INVENTION: OF PROSTATE CANCER THERAPIES AND THE DIAGNOSIS (
FILE REFERENCE: 07334-074001

CURRENT APPLICATION NUMBER: US/09/220,132

CURRENT FILING DATE: 1998-12-23

PRIOR APPLICATION NUMBER: US 60/079,303

PRIOR APPLICATION NUMBER: US 60/079,303

PRIOR APPLICATION NUMBER: US 60/068,821

PRIOR APPLICATION NUMBER: US 60/068,821

PRIOR APPLICATION NUMBER: US 60/068,821

PRIOR PRIOR PRIOR PRIOR DATE: 1997-12-24

NUMBER OF SECOND NOTE: 1997-12-24
TYPE: DNA ORGANISM: Homo sapiens FEATURE:
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LOCATION: 40..195
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                                                                                           for Windows
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97.8%;
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Pred. No. 6.5e-110;
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CANCER

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APPLICANT: Duclert, Aymeric
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS
FILE REFERENCE: 31.US3.CIP
CURRENT FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 09/191,997
PRIOR FILING DATE: 1998-11-13
PRIOR APPLICATION NUMBER: 60/066,677
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/069,957
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/096,573
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,273
PRIOR FILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 229
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GENERAL INFORMATION:
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Best Local Similarity
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LOCATION: (1)...(526)
OTHER INFORMATION: n =
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TNAAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCGTGTATTCTT
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99.0%;
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Pred. No. 1.2e-109;
0; Mismatches 3;
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Bougueleret, L.
APPLICANT: Jobert, S.
APPLICANT: Jobert, S.
TITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides
FILE REFERENCE: GENSET.050CP3
CURRENT APPLICATION NUMBER: US/09/599,360B
CURRENT FILING DATE: 2000-06-21
PRIOR APPLICATION NUMBER: 60/113,686
PRIOR APPLICATION NUMBER: 60/113,686
PRIOR APPLICATION NUMBER: 60/141,032
PRIOR APPLICATION NUMBER: 60/141,032
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 123
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm

TYPE: DNA

LENGTH:

Sequence 70, Application US/09599360B Patent No. 6548633 GENERAL INFORMATION:

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; NAME/KEY: polyA site
; LOCATION: 402..413
US-09-663-600A-147
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US-09-599-360B-70
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NAME/KEY: sig_peptide
LOCATION: 46.189
OTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 4.09999990462257
OTHER INFORMATION: seq VFMLIVSVLALIP/ET
NAME/KEY: polyA signal
LOCATION: 377.382
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SEQ ID NO 147
LENGTH: 413
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Best Local Similarity
Matches 412; Conserv
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                                                             CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAGAGTTTTGTAATTTTATATTACTTTT
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                                                                                                                      CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAGCTTTTGTAATTTTATATTACTTTT
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Pred. No. 7.7e-96;
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RESULT 8
US-09-385-982-95
; Sequence 95, Application
; Patent No. 6263334
; GENERAL INFORMATION:
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Best Local Similarity
Matches 504; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: sig_peptide
LOCATION: 140..442
OTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 4.10
OTHER INFORMATION: seq VFMLIVSVLALIP/ET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: polyA signal LOCATION: 630..635
NAME/KEY: polyA site LOCATION: 655..669
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                                                                                                                                                                                TTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCA
                                                                                                                                                                                                                                                             TTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCA 395
                                                                                                                                                                                                                                                                                                 CTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCAC
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                                                                                       AAAAAAA 669
                                                                                                                AAAAAAA 522
                                                                                                                                           TTTTTATCATCGCACAAGCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCG 302
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                           US/09385982
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Pred. No. 6.2e-73;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4;
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APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GE
TITLE OF INVENTION: PRODUCTS: II
FILE REFERENCE: CCDNA-260XX
CURRENT FILING DATE: 1999-0528,111
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-01-27
EARLIER FILING DATE: 1999-08-08
SARLIER FILING DATE: 1999-08-08
EARLIER FILING DATE: 1999-08-08
SARLIER FILING DATE: 1999-01-27
EARLIER FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 544
SOFTWARE: FASEUSEQ for Windows Version 3.0
SEQ ID NO 95
                                                                                                                                                                                                                                                                                                                                           RESULT 9
US-08-545-196B-10
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                                                                                                                                                                                                                                                                                       Sequence 10, Application US/08545196B Patent No. 6080577 GENERAL INFORMATION:
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Best Local Similarity
Matches 297; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(392)
OTHER INFORMATION: n = A,T,C o:
STATE: VA
STATE: VA
COUNTRY: USA
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                 NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: BIRCH, STEWART,
STREET: PO BOX 747
                                                                                                                                                                                                   APPLICANT: MELKI, JUDITH
APPLICANT: MUNNICH, ARNOLD
TITLE OF INVENTION: SURVIVAL MOTOR NEURON (SMN)
TITLE OF INVENTION: FOR SPINAL MUSCULAR ATROPHY
NUMBER OF SEQUENCES: 65
                                                                                                                               STREET: PO BOX 747
CITY: FALLS CHURCH
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95.5%;
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Pred. No. 2.4e-60;
0; Mismatches 12
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                                                                                                                                                                     KOLASCH
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US-08-545-196B-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: MELKI, JUDITH
APPLICANT: MUNNICH, ARNOLD
TITLE OF INVENTION: SURVIVAL MOTOR NEURON (SMN) GENE: A GENE
TITLE OF INVENTION: FOR SPINAL MUSCULAR ATROPHY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 205-8050 INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS:
                                                            TELEFAX: (703) 205-80
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 32,350
REFERENCE/DOCKET NUMBER: 21:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 19-OCT-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CUREENT APPLICATION DATA:
APPLICATION NUMBER: US/08/545,196B
                                                                                                      NAME: FARACI, C. J.
REGISTRATION NUMBER: 32,350
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 205-8000
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: FARACI, C. J.
                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 19-OCT CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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STRANDEDNESS:
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                               ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                               22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FALLS CHURCH
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                               1582 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: BIRCH, STEWART, KOLASCH AND BIRCH, LLP
PO BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1582 base pairs
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71.7%;
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Pred. No. 0.0012;
0; Mismatches 26;
                                                                                                                                                 2121-110P
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US-08-545-196B-12
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                                                                                                                            US-08-865-297-5
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                                                               Matches
                                                                            Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Prasad
APPLICANT: Schlos
                                                                                                                                                                                                                                                                                            TELEFAX: (617)227-5941 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: DF
                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: The Siva Genes, NO. TITLE OF INVENTION: CD27-Mediated Apoptosis
                                                                                                                                                                                           MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9.4%;
Local Similarity 71.7%;
les 66; Conservative
                                                                                                                                             NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                          NAME: Silveri, Jean M. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 29-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Boston
                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                             ENGTH:
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 628 CAGCACAAGATGTTCACACTAAAGAGAGAGAGGTGGCTTTTTATATGTTATGTTTTATA 687
                              413 CAGAAAAAGCCTGTGCATGAAAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTT
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                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: LAHIVE & COCKFIELD, LLP 60 State Street, Suite 510
                                                                                                                                                                                                                                                          770 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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88..612
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                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                              (617)227-7400
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                                                                                                                                                                                                                          single
                                                                              8.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The Siva Genes, No. 6010853el Genes Involved in
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Pred. No. 0.0012; 
0; Mismatches 2
                                                                          Score 46.8; DB 3; Length 770; Pred. No. 0.0072;
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RESULT 13
5194596-16
5194596-16
5194596
FARENT NO. 5194596
FAPPLICANT: TISCHER, EDMUND G.;ABRAHAM, JUDITH A.;FIDDES, JOH
COMITCHELL, RICHARD L.
TITLE OF INVENTION: PRODUCTION OF VASCULAR ENDOTHELIAL CELL
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GROWTH FACTOR
NUMBER OF SEQUENCES: 32
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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Patent No. 6307019
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212)391-0526
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Constar
APPLICANT: Zeng, 1
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NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                      438 AAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCT
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1185 Avenue of the Americas
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                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA (genomic)
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                                                                                                          EDMUND G.; ABRAHAM, JUDITH A.; FIDDES, JOHN
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   US/07/450,883
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                  CURRENT FILING DATE: 1998-12-04
EARLIER APPLICATION NUMBER: PCT/US98/11422
EARLIER FILING DATE: 1998-06-04
EARLIER APPLICATION NUMBER: 60/048,885
EARLIER FILING DATE: 1997-06-06
EARLIER APPLICATION NUMBER: 60/049,375
EARLIER FILING DATE: 1997-06-06
EARLIER APPLICATION NUMBER: 60/048,881
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US-09-205-258-64
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                                                                                                                                                                                                                                                  Sequence 64, Application US/09205258 Patent No. 6525174
                                                                                                                                                                                                                                   GENERAL
                                                                                                                                                                                               ENERAL INFORMATION:
APPLICANT: Young et al.
TITLE OF INVENTION: 207 Human Secreted Proteins
                                                                                                                                                              FILE REFERENCE: PZ007P1
CURRENT APPLICATION NUMBER: US/09/205,258
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FILING DATE: 1997-06-06

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Query Match
Best Local Similarity
Matches 69; Conserv
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                                                                                                                                                                       5219739-16
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; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 387,9
; FILING DATE: 27-JUL-1989
; SEQ ID NO:16:
                                                                                                                                                                                                                                                                                                                                                         JOHN C.; MITCHELL, RICHARD L.

TITLE OF INVENTION: DNA SEQUENCES ENCODING BVEGF120 AND HUMAN
HVEGF 121 AND METHODS FOR THE PRODUCTION OF BOVINE AND HUMAN
                                                                                                                                                                                                                                                                                                                                             VAASCULAR ENDOTHELIAL CELL GROWTH FACTORS, BVEGF120 AND HVEGF121
                                                                                                                                                                                                  SEQ ID NO:16:
                                                                                                                Matches
                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                        FILING DATE: 27-JUL-1990 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                              Local
                                                                                                                                                                                      LENGTH: 961
                                                                                                                                                                                                                                           APPLICATION NUMBER: 450,883 FILING DATE: 14-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 961
                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                 APPLICATION NUMBER: 387,545 FILING DATE: 27-JUL-1989
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
426 TGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTTGATACTAAGTATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         426 TGCATGAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATT
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                                                                                                                              Similarity
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                                                                                                               Conservative
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                                                                                                               Score 45; DB 6;
Pred. No. 0.023;
0; Mismatches
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Pred. No. 0.
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60/048,880

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EARLIER
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                  APPLICATION NUMBER: 60/092,921
FILING DATE: 1998-07-15
APPLICATION NUMBER: 60/094,657
FILING DATE: 1998-07-30
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APPLICATION NUMBER: 60/048,964
FILING DATE: 1997-06-06
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APPLICATION NUMBER: 60/048,963
EILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,877
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APPLICATION NUMBER: 60/048,898
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,962
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APPLICATION NUMBER: 60/048,897
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FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,892
FILING DATE: 1997-06-06
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FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,900
                                                                                                                      APPLICATION NUMBER: 60/070,923
FILING DATE: 1997-12-18
                                                                                                                                                                        FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,878
FILING DATE: 1997-06-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/048,883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/048,974
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APPLICATION NUMBER: 60/048,949
FILING DATE: 1997-06-06
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FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,917
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APPLICATION NUMBER: 60/048,875
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APPLICATION NUMBER: 60/049,373
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FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/049,019
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APPLICATION NUMBER: 60/048,899
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SEQ ID
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I LOCATION: (565)
I OTHER INFORMATION: n
FEATURE:
I FEATURE:
LOCATION: (566)
COTHER INFORMATION: n
US-09-205-258-64
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SEQ ID NO 64
LENGTH: 588
                                                                                                           Query Match
Best Local Similarity
Matches 59; Conserv
                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens FEATURE: NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                             TYPE: DNA
500
                          512 АААААААААААААААААААА 534
                                                     440 TTTTTTATTCCTGTGGGGCTCACCCCAAAGTATTAAAAGTAGCTTTGTAATTCCAAAAA
                                                                          TTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCAAAA
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                                                                                                           Score 44.6; DB 4;
Pred. No. 0.024;
0; Mismatches 24;
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                                                                                                                                       Length 588;
                                                                                                              Indels
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499 511 0

Search completed: January 31, 2004, 04:11:09 Job time: 79 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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seq length: 2000000000
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534
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nn2 6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
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6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
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Pred. No. and is No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.

SUMMARIES

15	14	13	12	11	10	9	œ	7	σ	σ	4.	ω	N	1	Result No.
335	368.4	404	411.4	425	460.2	466.8	478.6	478.6	486.6	486.6	497.2	507	509.4	534	Score
62.7	69.0	75.7	77.0	79.6	86.2	87.4	89.6	89.6	91.1	91.1	93.1	94.9	95.4	100.0	Query
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10	12	11	13	11	Ε	13	13	片	13	11	15	12	13	9	BB
US-09-978-295A-189	US-10-242-535A-7940	US-09-918-995-6534	US-10-319-763-147	US-09-822-846-397	US-09-918-995-19489	US-10-319-763-53	US-09-882-171-252	US-09-809-391-252	US-09-882-171-98	US-09-809-391-98	US-10-106-698-1927	US-10-291-172-13	US-10-119-428-14	US-09-801-115-1	ID
Sequence 189, App	Sequence 7940, Ap	Sequence 6534, Ap	Sequence 147, App	Sequence 397, App	Sequence 19489, A	Sequence 53, Appl	Sequence 252, App	Sequence 252, App	Sequence 98, Appl	Sequence 98, Appl	Sequence 1927, Ap	Sequence 13, Appl	Sequence 14, Appl	Sequence 1, Appli	Description

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16
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ALIGNMENTS

RESULT 1

US-09-801-115-1

Sequence 1, Application US/09801115
Patent No. US20020001828A1
GENERAL INFORMATION:

APPLICANT:

Ma, D.
Han, W.
Zhang, Y.
Song, Q. Song, Di, (

APPLICANT:

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TITLE OF INVENTION: CHEMOKINE LIKE FACTOR (CKLF) WITH CHEMOTACTIC AND TITLE OF INVENTION: STIMULATING ACTIVITIES
FILE REFERENCE: 10776-003-999
CURRENT APPLICATION NUMBER: US/09/801,115
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: PCT/CN00/00026
PRIOR PILING DATE: 2000-02-15
PRIOR PILING DATE: 2900-02-15
PRIOR APPLICATION NUMBER: 99107284.7
PRIOR FILING DATE: 1999-05-14
NUMBER OF SQ ID NOS: 8
SEQ ID NO 1
LENGTH: 534
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                     Query Match
Best Local Similarity
                                                                                                   Matches 534;
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 Di, C.
Huang, J.
Tang, J.
Chen, G.
                                                                     GTTCCCAATCTGAAGTGAAGCCGAGCTGGGCGAGAAGTAGGGGAAGGCCGTGCTCCGCCG
                                                                                                     Conservative
                                                                                                                   100.0%; Score 534;
100.0%; Pred. No. 8
                                                                                                   0; Mismatches
                                                                                                                     8.7e-123;
                                                                                                                                     Length 534;
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; NAME/KEY: CDS
; LOCATION: (236)..(535)
US-10-119-428-14
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CURRENT FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 09/596,193
PRIOR FILING DATE: 2000-06-17
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 55
SOFTWARE: Dt FL_genes Version 1.0
SEO ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Xu, Chongjun
APPLICANT: Mehrman, Tom
APPLICANT: Ren, Feiyan
APPLICANT: An, Yunqing
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Yang, Yonghong
APPLICANT: Yang, Yonghong
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: No. US20030165881A1el Nucleic Aci
TITLE REFERENCE: 789CIP2
FILE REFERENCE: 789CIP2
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US-10-119-428-14
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APPLICANT: Liu, Chenghua
APPLICANT: Asundi, Vinod
APPLICANT: Xu, Chongjun
APPLICANT: Wehrman, Tom
APPLICANT: Wehrman, Tom
APPLICANT: Ren, Felyan
APPLICANT: May Yunqing
APPLICANT: May Yunqing
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Oing A.
Query Match
Best Local Similarity
Matches 510; Conserv
                                                                                                                                                                                                                               SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/10119428
Publication No. US20030165881A1
                                                                                                                                           LENGTH: 817
TYPE: DNA
ORGANISM: Homo &
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TCGCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAA 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGAAAGTGCTGCTGCTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCCTGTGCATGAAAAAAAAAGATTTTGTAATTTTATATTTACTTTTAGTTTGATACTAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCCTGTGCATGAAAAAAAAGATTTTGTAATTTTATATTTACTTTTTAGTTTGATACTAA 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGC
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95.4%; ilarity 99.8%; Conservative
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Score 509.4; DB 13; Pred. No. 1.4e-116; 0; Mismatches 1;
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                                    Length
                                           817;
0;
  Gaps
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                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/291,172
CURRENT FILING DATE: 2000-11-08
PRIOR APPLICATION NUMBER: 09/63,267
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-09-65,363
PRIOR FILING DATE: 2000-09-66,847
PRIOR FILING DATE: 2000-09-14
PRIOR APPLICATION NUMBER: 09/616,847
PRIOR APPLICATION NUMBER: 09/56,193
PRIOR FILING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/574,954
PRIOR APPLICATION NUMBER: 09/579,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 752
SEQ ID NO 13
LENGTH: 655
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US-10-291-172-13
; Sequence 13, Application US/10291172
; Publication No. US20030228584A1
; GENERAL INFORMATION:
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                                                                                   US-10-291-172-13
Query Match
Best Local Similarity
Matches 507; Conserv
                                                                                                LENGTH: 655

TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (134)..(430)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hyseq, Inc
TITLE OF INVENTION: No. US20030228584A1el Nucleic
FILE REFERENCE: 21272-045
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  ilarity 100.0%;
Conservative (
94.9%; Score 507; DB 12;
100.0%; Pred. No. 5.1e-116;
live 0; Mismatches 0;
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                                           Length 655;
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RESULT 4
US-10-106-698-1927
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, NAME/KEY: misc feature

; LOCATION: (514)..(514)

; OTHER INFORMATION: n equals a,t,g, or

US-10-106-698-1927
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                                                                                                                                                                                                                                                                           FILE REFERENCE: PA005P1
CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR FILING DATE: 1999-09-29
PRIOR FILING DATE: 1999-09-29
PRIOR APPLICATION NUMBER: US 60/163,280
PRIOR APPLICATION NUMBER: US 60/163,280
PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 3.0
SEQ ID NO 1927
LENGTH: 558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1927, Application US/10106698 Publication No. US20030109690A1
                                                                          Query Match
Best Local 9
                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ruben et al.
TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypeptide
                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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 TGAAGYSGGAGCTGGCGAGAAGTAGGGGGAGGCGGTGCTCCGCCGCGGTGGCGGTTGCTA
                   TGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCCTCCGCCGCGGTGGCGGTTGCTA 75
                                                          93.1%;
nilarity 98.4%;
Conservative
                                                          ; Score 497.2; ; Pred. No. 1.3e 2; Mismatches
                                                       .2; DB 15;
.1.3e-113;
                                                               Indels
                                                                                         Length
                                                                                           558;
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; TYPE: DNA
; ORGANISM: Homo s
US-09-809-391-98
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US-09-809-391-98
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NUMBER OF SEQ ID NOS: 761
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Best Local Similarity
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TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16
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APPLICATION NUMBER: 60/056,845
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APPLICATION NUMBER: 60/056,636
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APPLICATION NUMBER: 60/056,862
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CURRENT FILING DATE: 2001-03-16
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TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
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Pred. No. 5.2
                                                    Score 478.6; DB 11
Pred. No. 5.1e-109;
2; Mismatches 11;
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PRIOR APPLICATION NUMBER: 60/040,334 PRIOR APPLICATION NUMBER: 60/040,336 PRIOR APPLICATION NUMBER: 60/040,336 PRIOR APPLICATION NUMBER: 60/040,163 PRIOR FILLING DATE: 1997-03-07 PRIOR PRIOR PRILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,615 PRIOR APPLICATION NUMBER: 60/047,597 PRIOR PRILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,597 PRIOR PRILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,633 PRIOR APPLICATION NUMBER: 60/047,633 PRIOR APPLICATION NUMBER: 60/047,633 PRIOR APPLICATION NUMBER: 60/047,633	2-171-252 10 252 10 252, Application US/098 11 INFORMATION: 2 NAT: Ruben et al. 2 INFORMATION: 3 INFORMATION: 3 INFORMATION: 186 Human S REFERENCE: PZ002P2 17 FILING DATE: 2001-05-1 4 APPLICATION NUMBER: 09/80 FILING DATE: 1998-09-08 APPLICATION NUMBER: PCT/U APPLICATION NUMBER: PCT/U FILING DATE: 1998-03-06 APPLICATION NUMBER: 60/04 FILING DATE: 1997-03-07 APPLICATION NUMBER: 60/04 FILING DATE: 1997-03-07 APPLICATION NUMBER: 60/03 FILING DATE: 1997-03-07	Qy 97 GGCAGCCAGCTGAGAAAGAGTTGAGGGAAAGTGCTGCTGGTGCAGACGCGATGGA 156 Db 62 GGCAGCCAGCTGAGAAATAAAACATCACTGCTGCTGCTGCTGCAGACGCGATGGA 121 Qy 157 TAACGTGCAGCCGAAAATAAAACATCACTCGCTTCTGCTTCAGTGTGAAAGCCCACTGAA 121 Db 122 TAACGTGCAGCCGAAAATAAAACATCACTCCCCCTTCTGCTTCAGTGTGAAAGCCCACGTGAA 181 Qy 217 GATGCTGCAGCCGAAAATAAAACATCACTCACTGTAACAACAGTATTCCATGCTCATCGTATC 276 Db 182 GATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATC 241 Db 182 GATGCTGCGCCTGATACCAGAAACCACACATTGACAACAGTATTCATGCTCATCGTATC 241 Qy 337 TGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAAGGGGTGTTTGCACT 301 Qy 337 TGTGACAGCAGTATGCTGTCTTGCCGAACATTGACAGTTGGTGGAAGGGTGTTTGCACT 396 Db 302 TGTGACAGCAGTATGCTGTTCTTGCCGAACGTTTTACCGGAAGCTTCGTTCA 361 Qy 397 TCCCAGGGGTCCTTACCAGAAAAAGCCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCA 361 Qy 397 TCCCAGGGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAA
PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION UNWBER: 60/048,97 PRIOR FILING DATE: 1997-06-06 PRIOR APPLICATION UNMER: 60/056,88 PRIOR FILING DATE: 1997-08-22 PRIOR APPLICATION NUMBER: 60/056,87 PRIOR TILING DATE: 1997-08-22 PRIOR FILING DATE: 1997-08-23 PRIOR PILING DATE: 1997-08-26 PRIOR APPLICATION NUMBER: 60/056,63 PRIOR APPLICATION NUMBER: 60/056,87	PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,60 PRIOR APPLICATION NUMBER: 60/043,58 PRIOR APPLICATION NUMBER: 60/043,58 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,31 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,31 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,31 PRIOR APPLICATION NUMBER: 60/043,31 PRIOR APPLICATION NUMBER: 60/043,31 PRIOR APPLICATION NUMBER: 60/043,67 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,67 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,67 PRIOR APPLICATION NUMBER: 60/043,67 PRIOR FILING DATE: 1997-04-11 PRIOR APPLICATION NUMBER: 60/043,31	PRIOR APPLICATION NUMBER: 60/047,58 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,61 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,61 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,50 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR FILING DATE: 1997-05-23 PRIOR PRILING DATE: 1997-05-23 PRIOR PRILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,49 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,49 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR APPLICATION NUMBER: 60/047,59 PRIOR FILING DATE: 1997-05-23 PRIOR APPLICATION NUMBER: 60/047,59

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Sequence 53, Application US/10319763
Publication No. US20030144490A1
GENERAL INFORMATION:
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Best Local
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                                          APPLICANT: Duclert, Aymeric
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: EXTENDED COMAS FOR SE
FILE REFERENCE: G-031.US04.DIV
CURRENT APPLICATION NUMBER: US/10/319,763
CURRENT FILING DATE: 2002-12-10
                                                                                                                      APPLICANT: Dumas Milne Edwards,
APPLICANT: Duclert, Aymeric
PRIOR APPLICATION NUMBER: 60/066,677
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/069,957
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PR APPLICATION NUMBER: 60/048,964

DR FILING DATE: 1997-06-06

PR APPLICATION NUMBER: 60/057,650

DR FILING DATE: 1997-09-05

DR APPLICATION NUMBER: 60/056,884

DR FILING DATE: 1997-08-22

DR APPLICATION NUMBER: 60/057,669

DR FILING DATE: 1997-09-05

DR FILING DATE: 1997-09-05
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FILING DATE:
APPLICATION N
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APPLICATION NUMBER: 60/056,887
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,908
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                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAA
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Pred. No. 5.1e-109;
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OR APPLICATION NUMBER: 60/056,864
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,631
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,845
OR APPLICATION NUMBER: 60/056,892
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/057,761
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,599
OR APPLICATION NUMBER: 60/047,598
OR APPLICATION NUMBER: 60/047,588
OR APPLICATION NUMBER: 60/047,585
OR APPLICATION NUMBER: 60/047,586
OR APPLICATION NUMBER: 60/047,580
OR APPLICATION NUMBER: 60/047,590
OR FILING DATE: 1997-05-23

OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,880
OR PILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,894
OR PILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,911
OR PILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,36
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,874
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,874
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,910
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,910
OR FILING DATE: 1997-08-22

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OR FILING DATE: 1997-08-22

REAPPLICATION NUMBER: 60/056,903

REFILING DATE: 1997-08-22

REAPPLICATION NUMBER: 60/056,888

OR FILING DATE: 1997-08-22

REAPPLICATION NUMBER: 60/056,879

APPLICATION NUMBER: 60/056,872
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,882
FILING DATE: 1997-08-22
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,637

1997-08-22

PRIOR PRIOR

FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,576
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/047,501
FILING DATE: 1997-05-23

APPLICATION NUMBER: 60/047,589
FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,593
FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,614

FILING DATE:

1997-05-23

FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,578

OR APPLICATION NUMBER: 60/043,670
R FILING DATE: 1997-04-11
OR APPLICATION NUMBER: 60/056,632
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,664
OR FILING DATE: 1997-08-22
OR PILING DATE: 1997-08-22
OR PILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,876
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,881

APPLICATION N FILING DATE:

NUMBER:

60/056,909

1997-08-

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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR APPLICATION NUMBER: 60/086,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR FILING DATE: 1998-08-10
PRIOR PILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 229
SOFTWARE: Patent.pm
SEQ ID NO 53
LENGTH: 500
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: sig_peptide
LOCATION: 199..288
OTHER INFORMATION: secore 5.5999990463257
OTHER INFORMATION: seq IVSUALIPETTI/LT
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc feature
LOCATION: 197.-412
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
                                                                     COATION: 207..412
OTHER INFORMATION: b
OTHER INFORMATION: b
OTHER INFORMATION: 6
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OTHER INFORMATION: 1
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OTHER INFORMATION: 1
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OTHER INFORMATION: 6
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LOCATION: 61.195
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA455042
OTHER INFORMATION: est
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LOCATION: 425..488
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
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LOCATION: 61.195
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
NAME/KEY: misc_feature
LOCATION: 58..195
OTHER INFORMATION: homology
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LOCATION: 464..469
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LOCATION: 489..500
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JOCATION: 197._412
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id :AA455042
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id :W93646
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FEATURE:
NAME/KEY: misc feature
LOCATION: 197..412
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA516431
OTHER INFORMATION: est
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LOCATION: 197..338
OTHER INFORMATION: homol
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LOCATION: 425..469
OTHER INFORMATION: homo:
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LOCATION: 71..195
OTHER INFORMATION: homology
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OTHER INFORMATION: est
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LOCATION: 443..477
OTHER INFORMATION: homology
OTHER INFORMATION: id: W388
OTHER INFORMATION: est
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LOCATION: 197.324
OTHER INFORMATION: homology
OTHER INFORMATION: id: w38899
OTHER INFORMATION: est
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LOCATION: 52..195
OTHER INFORMATION: homology
OTHER INFORMATION: id :w38899
OTHER INFORMATION: est
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LOCATION: 90.195
OTHER INFORMATION: homology
OTHER INFORMATION: id: AA516431
OTHER INFORMATION: est
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OTHER INFORMATION: id :W93646
OTHER INFORMATION: est
NAME/KEY: misc_feature
LOCATION: 40..195
OTHER INFORMATION: homology
OTHER INFORMATION: id :W19506
                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: 339..401
OTHER INFORMATION: hom
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LOCATION: 425..488
OTHER INFORMATION: homology
OTHER INFORMATION: id ;AA516431
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LOCATION: 425..488
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id :W52820
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id:W52820
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                                                                                                  PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 19489
LENGTH: 512
Type: name
                                                                                                                                                                                                                                                                                                 Sequence 19489, Application US/09918995
Publication No. US20030073623A1
GENERAL INFORMATION:
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (1):..(512)
OTHER INFORMATION: n = A,T,C
-09-918-995-19489
                                                                                                                                                                                                  APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
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NAME/KEY: misc_feature
LOCATION: 9..10,12
OTHER INFORMATION: n=a,
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                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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Pred. No. 4.4e-106;
4; Mismatches 5;
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GENERAL INFORMATION.

APPLICANT: Jacobs, Kenneth

APPLICANT: McCoy, John M.

APPLICANT: LaVallie, Edward R.

APPLICANT: LaVallie, Edward R.
                            APPLICANT: Gulukota, Kamalakar
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILE REFERENCE: GIN 6400
CURRENT APPLICATION NUMBER: US/09/822,846
CURRENT APPLICATION NUMBER: 001-03-29
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR TILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 629
SOOTWARE: Patentin Ver. 2.0
SEQ ID NO 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 397, Application US/09822846 Publication No. US20030027139A1
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Best Local
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LENGTH: 43
TYPE: DNA
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                                                                                                                                                                                                                            Howes, Steven H.
Resnick, Richard J.
                                                                                                                                                                                                                                                            Fechtel, Kim
                                                                                                                                                                                                                                                                             Clark, Hilary
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Wong, Gordon G.
                                                                                                                                                                                                                                                                                                                         Agostino, Michael J.
Steininger II, Robert
Bowman, Michael R.
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Treacy, Maurice
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99.1%;
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Pred. No. 1.9e-104;
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ORGANISM: Homo sapiens

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RESULT 12
US-10-319-763-147
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CURRENT APPLICATION NUMBER: US/10/319,763
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: 60/066,677
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/069,957
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR FILING DATE: 1998-08-10
PRIOR FILING DATE: 1998-08-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 147, Application US/10319763 Publication No. US20030144490A1 GENERAL INFORMATION:
                                                                                                                                       NUMBER OF SEQ ID NOS:
SOFTWARE: Patent.pm
SEQ ID NO 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: EXTENDED CONAS FOR SECRETED PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dumas Milne Edwards, Jean-Baptiste
                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/099,273 PRIOR FILING DATE: 1998-09-04
NAME/KEY: sig_peptide
LOCATION: 46..189
OTHER INFORMATION: Von Heijne matrix
                                                         TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                     LENGTH: 413
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Pred. No. 1e-95;
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                                                                                                                                                                                                                                                      Sequence 6534, Application US/09918995
; Publication No. US20030073623A1
; GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
INTIE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
; CURRENT APPLICATION NUMBER: US/09/918,995
; CURRENT APPLICATION NUMBER: US/09/918,995
; CURRENT APPLICATION NUMBER: US/09/235,076
; PRIOR APPLICATION NUMBER: US/09/235,076
; PRIOR APPLICATION NUMBER: US/09/235,076
; PRIOR FILING DATE: 1999-01-20
; NUMBER OF SEQ ID NOS: 38054
; SOFTWARE: FASTSEQ for Windows Version 3.0
; SEQ ID NO 6534
; LENGTH: 422
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; ORGANISM: Homo sapiens
US-09-918-995-6534
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US-09-918-995-6534
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                                                                                                                                       Query Match
Best Local &
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Best Local
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NAME/KEY: polyA_site
LOCATION: 402..413
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OTHER INFORMATION: seq VFMLIVSVLALIP/ET
                                                                                                                                         Local Similarity
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hes 412; Conserv
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                                                                              122 GAAAGTGCTGCTGCTGCGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAAACAT
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    182 CGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAAC
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                                             GAAAGTGCTGCTGCGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACAT
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                                                                                                                       75.7%; So ilarity 100.0%; E Conservative 0;
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                                                                                                                       Score 404; DB 11;
; Pred. No. 1.7e-90;
0; Mismatches 0;
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Pred. No. 2.4e-92;
0; Mismatches 1
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377 413 317 353 257

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SOFTWARE: PatentIn version 3.2
SEQ ID NO 7940
LENGTH: 400
TYPE: DNA
ORGANISM: Human
FEATURE:
NAME/KEY: misc feature
LOCATION: (176)..(176)
OTHER INFORMATION: n is a, c, g
US-10-242-535A-7940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7940, Application US/10242535A Publication No. US20040013663A1
                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 58994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis FILE REFERENCE: 4231/2005
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TATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACC
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PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
PRIOR FILING DATE: 1997-11-13
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PRIOR APPLICATION NUMBER: 60/077450
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APPLICANT: Ashkenazi, Avi
APPLICANT: Baker Kevin P
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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2630P1C11
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Shelton, David L.
Stewart, Timothy A.
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Hillan, Kenneth J
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Grimaldi, J. Christopher
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Filvaroff, Ellen
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Query Match

62.7%; Score 335; DB 10; Length 663;

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                                          PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                                         AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                        GTGAAGATGCTGCGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
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Mammalia; Eutheria; Primates;
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AF096895
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Molecular cloning and characterization of chemokine-like factor 1 (CKLF1), a novel human cytokine with unique structure and potential chemotactic activity

Blochem. J. 357 (Pt 1), 127-135 (2001)
Science Center, Xueyuan Road 38, Beijing 100083, China Sequence update by submitter On Sep 7, 2000 this sequence version replaced gi:6288733.
                        Direct Submission
Submitted (07-SEP-2000) Immunology, Peking University Health
Science Center, Xueyuan Road 38, Beijing 100083, China
                                                                                                Direct Submission
Submitted (03-OCT-1998) Immunology,
Xue Yuan Rd., Beijing 100083, China
3 (bases 1 to 530)
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Han, W.L., Li, Y., Zhang, Y.M.,
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FLI_CDNA.
                                                                                    Han, W.L.
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ALIPETTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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AR275027
                                     Methods and compositions for the identification and assessment prostate cancer therapies and the diagnosis of prostate cancer Patent: US 6506607-A 164 14-JAN-2003;

Location/Qualifiers
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Shyjan, A.W.
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aLIPETTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
108 c 137 g 129 t
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/organism="unknown"
136 c 113 g
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/cell_line="U937"
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/protein_id="AAF06722.1"
/db_xref="GI:6288734"
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PARENT: JP 2001523453-A 47 27-NOV-2001;

GENSET

OS Homo sapiens (human)

PD 27-NOV-2001

PP 13-NOV-1998 JP 2000521191

PF 13-NOV-1998 US 60/066677,17-DEC-1997 US 60/069957 PR

13-NOV-1997 US 60/074121,13-APR-1998 US 60/099273 PI LYDIE

809-FEB-1998 US 60/074121,13-APR-1998 US 60/099273 PI LYDIE

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PC C12N15/09,C12N15/09,C07K14/47,C07K16/18,C12N1/15,C12N1/19, PC

C12N1/21,

CC C12N15/10,C12P21/02,C12Q1/68,C12N15/00,C12N15/00,C12N15/00

CC SECQ IVSYLALIPETTY/LT

CC homology

CC id:AA429945

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                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 500)
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JP 2001523453-A/47.
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AF057306.1 GI:6648618
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                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Direct Submission
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
102 c 125 g 12
/codon_start=1
product="transmembrane proteolipid"
protein_id="AAF21255.1"
/brotein_id="AAF21255.1"
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/translation="MDNVQPKIKHRPFCFSVKGHVKMLRLALTVTSMTFFIIAQAPEP
YIVITGFEVTVILFFILLYVLRLDRLMKWLFWPLLDIINSLVTTVFMLIVSVLALIPE
                                                                /gene="C32"
/note="down regulated upon
sodium butyrate"
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132. .590
                                                                                                                                                            /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/cell_line="AsPC-1"
                                                                                                                                       note="adenocarcinoma"
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Edwards,J.-B.D.M., Bougueleret,L. and
Complementary DNA's encoding proteins
Patent: US 6548633-A 70 15-APR-2003;
Location/Qualifiers
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AX061665
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Mammalia; Eutheria; Primates;
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                                                                                                                                                /note="Von Heijne matrix score 630. .635
                                                                                                                                                                                                                                                      /protein_id="CAC25025.1"
/db_xref="GI:12406790:
/db_xref="GI:12406790:
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YIVITGFEVTVILFFILLYVLRLDRLMKWLFWPLLDIINSLVTTVFMLIVSVLALIPE
                                                                                                                                                                                                                              TTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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/mol type="genomic DNA"
/db xref="taxon:9606"
140. .598
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KEYWORDS
SOURCE
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LOCUS
DEFINITION
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JOURNAL
                                                                                                                                              NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A. G.E. Consortium (LLNL)
DNA Sequencing by: Genome Sequence Centre,
BC Cancer Agency, Vancouver, BC, Canada
info@bcgsc.bc.ca
                  Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hejao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabbu, Parvaine Spacedi, Jacqueline Cheir, Parvaine Spacedi, Jacqueline Cheir, Parvaine Spacedi, Jacqueline Cheir, Parvaine Carrier, Martin Martin, Martin Martin, Mart
                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (12-MAR-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MGC
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BC004380.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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  Smailus, Michael Smith,
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IMAGE:3639550, mRNA,
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complete cds. PRI 12-JUL-2001

Lorraine Spence,

Jeff Stott,

Score: Percent Similarity Best Local Similar Query Match: DB:	e: 471.50 Matches: 99 ent Similarity: 65.13% Conservative: 0 Local Similarity: 65.13% Mismatches: 0 y Match: 92.81% Indels: 53 y Match: 6 Gaps: 1
US-09-801-115B-2	-115B-2 (1-99) x AX061665 (1-669)
γ	1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis '20
Вb	140 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC 199
Qy	21 VallysMetLeuArg 25
Вb	200 GTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGCACAA 259
δ	25 25
Db	260 GCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTTCATA 319
Qy	26Leuhap 27
Вb	320 CTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCTTGAT 379
ν,	28 IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle 47
Db .	380 ATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGCACTGATA 439
γ	48 ProGluThrThrLeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCys 67
Вb	440 CCAGAAACCACAACATTGACAGTTGGTGGAGGGGGTGTTTGCACTTGTGACAGCAGTATGC 499
δ	68 CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr 87
Db dd	500 TGTCTTGCCGACGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTAC 559
Qy	88 GlnLysLysProValHisGluLysLysGluValLeu 99
Вb	560 CAGAAAAAGCCTGTGCATGAAAAAAAAAGAAGTTTTG 595
RESULT 10	

Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.

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RESULT 11
AF135380
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DB:
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Best Local Similarity:
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    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     through the I.M.A.G.E. Consortium/LLNL at: http://image.lln
Series: IRAL Plate: 13 Row: f Column: 14
This clone was selected for full length sequencing because
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clone distribution: MGC clone distribution information can be found
                                                                                 GlnLysLysProValHisGluLysLysGluValLeu
                                                                                                                                                                                                           CCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VallysMetLeuArg------
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                                                         CAGAAAAAGCCTGTGCATGAAAAAAAAAAGAAGTTTTG 592
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TTTLTYQGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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/clone="MGC:10658 IMAGE:3639550"
/fissue_type="Pancreas, adenocar/
/clone_Tib="NIH_MGC_39"
/lab_host="DH10B-R"
/lab_mst="DH10B-R"
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Location/Qualifiers
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Submitted (17-MAR-1999) Immunology, Beijing Medical University,
Yuan Road, Beijing 100083, China
2 (bases 1 to 689)
                                                                                                                                                                                                                                                                                         1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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Han, W.L., Gu, W., Li, Y., Zhang, Y., Song, Q.,
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Mammalia; Eutheria; Primates;
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IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle
                                                                                                                                                                                       GTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGCACAA
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                                                   CTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCTTGAT
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/codon_start=1
/product="chemokine-like factor 2"
/product="chemokine-like factor 2"
/product="chemokine-like factor 2"
/product="chemokine-like factor 2"
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/db_xref="GI:6630854"
/translation="MNNVQPKIKHRPFCFSVKGHVKMLFLALTVTSMTFFIIAQAPEP
/translation="MNNVQPKIKHRPFCFSVKGHVKMLFLALTVTSMTFFIIAQAPEP
YIVITGFEVTVLIFFILLYVLRLDRLMKWLFWPLLDIINSLVTTVFMLIVSVLALIPE
TTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKFVHEKKEVL"
a 140 c 161 g 193 t
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/db_xref="taxon:9606"
/cell_line="U937"
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Road 38, Beijing
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Edwards, J. B.D.M., Duclair, B. and Jordan, J.Y.
Sequence tag and encoded human protein
Patent: JP 2001269182-A 138 02-OCT-2001;
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Homo sapiens
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ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyValPhe 60
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26-FEB-1999 US 60/122487
JEAN BAPUTIST DUMAS MILNE EDWARDS, EIMERIC DUCLAIR, JEAN YVES
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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RESULT 13
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                                                                                                         Prepared with primer pairs provided by Sandoz,
-- Washington University/Merck EST sequence.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Richard M. Myers
Stanford Human Genome Center (SHGC)
Stanford University School of Medicine
Department of Genetics, M-344, Stanford,
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G30204.1 GI:1593755
STS; STS sequence; p
                                                                                                                                                                                                           Buffer:
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primer B: GCCCTTATTTACCGGAAGCT
STS size: 77
PCR Profile:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: myers@shgc.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 4157259687
Fax: 4157259689
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Myers, R.M.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 485)
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                                                                                                                                                          Tris-HCl:
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KCl:
                         /map="16"
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Total Vol:
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PCR Cycles:
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 complement (126. .145)
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                                                           /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                 Science Center, Xueyuan Road 38, Beijing 100083, China
Sequence update by submitter
On Sep 7, 2000 this sequence version replaced gi:6625671
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                        Han, W.L.
                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (23-APR-1999) Immunology,
Yuan Road, Beijing 100083, China
2 (bases 1 to 593)
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             166
             ы
                         NPSGPYQKKPVHEKKEVL"
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                                                                                                                                           gene="CKLF4"
                                                                                                                                                                     note="synonym: UCK"
                                                                                                                                                                                                                cell_line="U937"
note="from PHA stimulated
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Beijing 100083, China
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                                                                                                                                     Submitted (07-SEP-2000) Immunology, Peking University Health Science Center, Xueyuan Road 38, Beijing 100083, China Sequence update by submitter On Sep 7, 2000 this sequence version replaced gi:6630855.
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Submitted (17-MAR-1999) Immunology,
Yuan Road, Beijing 100083, China
2 (bases 1 to 434)
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens chemokine-like alternatively spliced.
                                                                                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                Han, W.L
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Han, W.L., Gu, W., Li, Y., Zhang, Y., Song, Q.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                     /note="from PHA stimulated cells"
                                                                                                 organism="Homo sapiens"
|mol_type="mRNA"
             'note="UCK"
                          gene="CKLF3"
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Conservative:
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Indels:
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3 (CKLF3) mRNA,
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Percent Similarity:
Best Local Similarity:
Query Match:
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Search completed: January 31, 2004, 04:50:56 Job time : 1820 secs
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ORIGIN
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Pred. No.:
                                                                                                                                                                                                                                                                292
                                                                                                      232 GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGCCCCTTATTTACCGGAAGCTTCTG 291
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                                                  61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
                                                                                                                                                                                  41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         /gene="CKLF3"
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/note="UCK-3; alternatively spliced"
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/db_xref="G1.6630856"
/translation="MDNOPKIKHRPFCFSVKGHVKMLRLVFALVTAVCCLADGALIY
RKLLFNPSGPYQKKPVHEKKEVL"
a 88 c 116 g 103 t
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-QB=Published Applications NA -QFMT=fastap -SUFFIX=p2n.rnpb -MINMATCH=0.1
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-MODEL=frame+_p2n.model
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                    score greater than or equal
  Score
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Ygapop 10.0, Y
Fgapop 6.0, F
Delop 6.0, I
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Match Length DB
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// Cgn2_6/ptodata/1/pubpna/USO9_PUBCOMB.seq:*
// Cgn2_6/ptodata/1/pubpna/USO9_NEW_PUB.seq:*
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1321.652 Million cell updates/sec
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  Description
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508 100.0 413 13 US-10-319-763-147 Sequence 147, App 1508 100.0 432 11 US-09-818-995-6534 Sequence 374, App 508 100.0 504 11 US-09-809-391-98 Sequence 98, App 1508 100.0 506 11 US-09-809-391-298 Sequence 98, App 1508 100.0 506 11 US-09-809-391-252 Sequence 99, App 1508 100.0 506 11 US-09-809-391-252 Sequence 252, App 508 100.0 556 12 US-10-219-731-52 Sequence 1, App 1508 100.0 558 15 US-10-219-731-252 Sequence 1, App 1508 100.0 558 15 US-10-219-172-13 Sequence 1, App 1508 100.0 558 15 US-10-219-731-353 Sequence 1, App 1509 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559 100.0 559
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ALIGNMENTS

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RESULT 1

US-10-319-763-147

; Sequence 147, Application US/10319763

Publication No. US20030144490A1

; GENERAL INFORMATION:

APPLICANT: Dunas Milne Edwards, Jean-Baptiste

APPLICANT: Duclert, Aymeric

APPLICANT: Bougueleret, Lydie

TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS

FILE REFERENCE: G-031.US04.DIV

CURRENT APPLICATION NUMBER: US/10/319,763

; CURRENT APPLICATION NUMBER: 60/066,677

PRIOR APPLICATION NUMBER: 60/069,957

PRIOR FILING DATE: 1997-11-13

; PRIOR APPLICATION NUMBER: 60/074,121

PRIOR APPLICATION NUMBER: 60/081,563

PRIOR APPLICATION NUMBER: 60/096,116

PRIOR APPLICATION NUMBER: 60/096,116

PRIOR APPLICATION NUMBER: 60/096,116

PRIOR APPLICATION NUMBER: 60/096,273

PRIOR FILING DATE: 1998-08-10

; PRIOR FILING DATE: 1998-09-04
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Sequence 6534, Application US/09918995

| Sequence 6534, Application US/09918995
| Publication No. US20030073623A1
| GENERAL INFORMATION:
| APPLICANT: Hyseq, Inc.
| TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
| FILE REFERENCE: 20411-756
| CURRENT APPLICATION NUMBER: US/09/918,995
| CURRENT FILING DATE: 2001-07-30
| PRIOR APPLICATION NUMBER: US/09/235,076
| PRIOR APPLICATION NUMBER: US/09/235,076
| PRIOR APPLICATION NUMBER: US/09/235,076
| PRIOR FILING DATE: 1999-01-20
| NUMBER OF SEQ ID NOS: 38054
| SOFTMARE: FastSEQ for Windows Version 3.0
| SEQ ID NO 6534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 229

SOFTWARE: Patent.pm

SEQ ID NO 147

LENGTH: 413

TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: sig_peptide
LOCATION: 46..189
OTHER INFORMATION: SCOTE 4.0999990463257
OTHER INFORMATION: BEQ VFMLIVSVLALIP/ET
 Alignment Scores: Pred. No.:
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US-09-918-995-6534
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Best Local Similarity:
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                                              US-09-918-995-6534
                                                         LENGTH: 422
TYPE: DNA
ORGANISM: Homo sapiens
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NAME/KEY: polyA_site
LOCATION: 402..413
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Best Local Similarity:
Query Match:
DB:
Percent Similarity:
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Query Match:
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                                                                                                                                               ; LENGTH: 439
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-822-846-397
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APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: POLYMUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILE REFERENCE: GIN 6400
CURRENT APPLICATION NUMBER: US/09/822,846
CURRENT APPLICATION NUMBER: 001-03-29
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR TILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 629
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Jacobs, Kenneth APPLICANT: McCoy, John M.
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Wong, Gordon G.
Clark, Hilary
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LaVallie, Edward R.
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Howes, Steven H.
Resnick, Richard J.
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Steininger II, Robert
Bowman, Michael R.
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Treacy, Maurice
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Evans, Cheryl
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US-09-809-391-98
RESULT 5
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Publication No. US20030049618A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186
FILE REFERENCE: PZ002P2
CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 761
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
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                                                PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
                                                                                                             AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                          GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
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US-09-882-171-98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2001-03-16 PRIOR APPLICATION NUMBER: 09/149,476
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APPLICATION NUMBER: PCT/US98/04493
APPLICATION NUMBER: 60/047,612
FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,632
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APPLICATION NUMBER: 60/047,492
FILING DATE: 1997-05-23
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APPLICATION NUMBER: 60/047,615
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APPLICATION NUMBER: 60/047,600
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APPLICATION NUMBER: 60/040,163
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APPLICATION NUMBER: 60/040,333
FILING DATE: 1997-03-07
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                                                                    FILING DATE:
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FILING DATE: 1997-05-23
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APPLICATION NUMBER: 60/047,502
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R FILING DATE: 1997-04.
R APPLICATION NUMBER: R AFFLING DATE: 1997-04.
R FILING DATE: 1997-04.
R FILING DATE: 1997-04.

NUMBER: 60 1997-04-1

60/043,314

1997-04-

60/043,580 60/043,568

DR APPLICATION NUMBER: 60/043,569

RFILING DATE: 1997-04-11

RR APPLICATION NUMBER: 60/043,311

RR FILING DATE: 1997-04-11

RR APPLICATION NUMBER: 60/043,671

RR FILING DATE: 1997-04-11

RR APPLICATION NUMBER: 60/043,674

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OR FILLING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,664
OR FILLING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,876
OR APPLICATION NUMBER: 60/056,881
OR APPLICATION NUMBER: 60/056,881
OR APPLICATION NUMBER: 60/056,909
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,875
OR APPLICATION NUMBER: 60/056,875
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,862
OR APPLICATION NUMBER: 60/056,862
OR APPLICATION NUMBER: 60/056,862
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OR APPLICATION NUMBER: 60/056,968
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OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,630
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OR APPLICATION NUMBER: 60/056,662
OR PILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,662
OR APPLICATION NUMBER: 60/056,872
OR APPLICATION NUMBER: 60/056,872
OR FILING DATE: 1997-08-22
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OR APPLICATION NUMBER: 60/056,882
OR APPLICATION NUMBER: 60/056,903
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DR APPLICATION NUMBER: 60/056,889

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OR APPLICATION NUMBER: 60/056,864
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Sequence 252, Application US/09882171
Publication No. US20030175858A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
CURRENT APPLICATION NUMBER: US/09/882,171
CURRENT FILING DATE: 2001-06-18
PRIOR APPLICATION NUMBER: 09/809,391
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TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16
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OR APPLICATION NUMBER: 60/047,601
OR FILING DATE: 1997-05-23
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OR APPLICATION NUMBER: 60/043,568
OR APPLICATION NUMBER: 60/043,314
OR APPLICATION NUMBER: 60/043,314
OR FILING DATE: 1997-04-11
OR FILING DATE: 1997-04-11 OR FILING DATE: 1997-05-23
OR APPLICATION NUMBER: 60/047,613
OR PRILING DATE: 1997-05-23
OR APPLICATION NUMBER: 60/047,582
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FILING DATE: 1998-09-08 APPLICATION NUMBER: 60/043,569 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,311 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/047,632 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/040,162 1997-03-07 1997-05-23 1997-03-07 1997-03-07 1997-03-07 1997-05-23 1997-05-23 1997-03-07 1997-03-07 60/047,597 60/040,163 60/038,621

DR APPLICATION NUMBER: 60/043,671
DR FILING DATE: 1997-04-11
DR APPLICATION NUMBER: 60/043,674
DR FILING DATE: 1997-04-11
DR APPLICATION NUMBER: 60/043,669
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OR FILING DATE: 1997-04-11
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                                                                                                                          GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
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                                          GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGCCCCTTATTTACCGGAAGCTTCTG
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                                                                                                                                                                                                                                                                                                                                                                        (1-99) x US-09-882-171-252 (1-506)
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Matches:
Conservative:
Mismatches:
Indels:
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PRIOR PRIOR

OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,864
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,631
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,845
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,892
OR APPLICATION NUMBER: 60/057,761
OR APPLICATION NUMBER: 60/047,595
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/047,595
OR FILING DATE: 1997-05-23
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,595
OR APPLICATION NUMBER: 60/047,586
OR APPLICATION NUMBER: 60/047,586
OR APPLICATION NUMBER: 60/047,586
OR APPLICATION NUMBER: 60/047,586
OR APPLICATION NUMBER: 60/047,585

OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,879
OR PELING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,880
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,894
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,911
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,636
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,874
OR APPLICATION NUMBER: 60/056,874

FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,903
FILING DATE: 1997-08-25
APPLICATION NUMBER: 60/056,888

FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,637

OR APPLICATION NUMBER: 60/043,315
OR FILLING DATE: 1997-04-11
OR APPLICATION NUMBER: 60/056,886
OR APPLICATION NUMBER: 60/056,886
OR APPLICATION NUMBER: 60/056,870
OR APPLICATION NUMBER: 60/056,877
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,897
OR APPLICATION NUMBER: 60/056,897
OR APPLICATION NUMBER: 60/056,897
OR APPLICATION NUMBER: 60/056,897
OR APPLICATION NUMBER: 60/056,630
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,630
OR FILING DATE: 1997-08-22
OR APPLICATION NUMBER: 60/056,630
OR APPLICATION NUMBER: 60/056,878
OR APPLICATION NUMBER: 60/056,662
OR APPLICATION NUMBER: 60/056,662
OR APPLICATION NUMBER: 60/056,672
OR APPLICATION NUMBER: 60/056,872

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Percent Similarity:
Best Local Similarity:
Query Match:
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; LENGTH: 534
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-801-115-1
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APPLICANT: Han, W.
APPLICANT: Song, Y.
APPLICANT: Song, Q.
APPLICANT: Di, C.
APPLICANT: Huang, J.
APPLICANT: Tang, J.
APPLICANT: Chen, G.
TITLE OF INVENTION: CHEMOKINE LIKE FACTOR(CKL
TITLE OF INVENTION: STIMULATING ACTIVITIES
FILE REFERENCE: 10776-003-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-801-115-1
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                                                                                                                      GENERAL INFORMATION:
                                                                                                                                           Sequence 1927, Application US/10106698 Publication No. US20030109690A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/801,115
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: PCT/CN00/00026
PRIOR FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 99107284.7
PRIOR FILING DATE: 1999-05-14
NUMBER: 1999-05-14
APPLICANT: Ruben et al.
TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypeptide
FILE REFERENCE: PA005P1
CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                       41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 VallysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
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                                                                                                                                                                                                                                                                                PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
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FILE REFERENCE: 212772-045
CURRENT APPLICATION NUMBER: US/10/291,172
CURRENT FILING DATE: 2000-11-08
PRIOR APPLICATION NUMBER: 09/633,267
PRIOR APPLICATION NUMBER: 09/653,363
PRIOR FILING DATE: 2000-09-19
PRIOR APPLICATION NUMBER: 09/616,847
PRIOR APPLICATION NUMBER: 09/616,847
PRIOR APPLICATION NUMBER: 09/56,193
PRIOR FILING DATE: 2000-07-14
PRIOR FILING DATE: 2000-06-17
PRIOR PRIOR DATE: 2000-06-17
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR FILING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR PILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 752
SEQ ID NO 13
LENGTH: 655
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PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
SOFTWARE: PatentIn Ver. 3.0
SEQ ID NO 1927
SEQ ID NO 1927
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (514)...(514)
OTHER INFORMATION: n equals a,t,g, or c
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DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-106-698-1927
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/10291172
Publication No. US20030228584A1
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Hyseq, Inc
TITLE OF INVENTION: No. US20030228584A1el Nucleic Acids
FILE REFERENCE: 21272-045
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PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR FILING DATE: 1999-09-29
TYPE: DNA
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Matches:
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Gaps:
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206 20

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APPLICANT: Ren, Felyan
APPLICANT: Ma, Yunqing
APPLICANT: Zhou, Ping
APPLICANT: Zhao, Qing A.
APPLICANT: Zhao, Qing A.
APPLICANT: Yang, Yonghong
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: No. US20030165881A1el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 789CIP2
CURRENT APPLICATION NUMBER: US/10/119,428
CURRENT FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 09/596,193
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/574,454
PRIOR APPLICATION NUMBER: 09/579,705
PRIOR APPLICATION NUMBER: 09/519,705
PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 55
SOFTWARE: pt_FL_genes Version 1.0
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (236)..(535)
US-10-119-428-14
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Tang, Y
APPLICANT: Liu, G
APPLICANT: Asundir
APPLICANT: Xu, Ct
APPLICANT: Wehrma
APPLICANT: Ren, T
APPLICANT: Ren, T
APPLICANT: Ren, T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: CDS
LOCATION: (134)..(430)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81 PheAsnProSerGlyProTyrGlnLysLysBroValHisGluLysLysGluValLeu 99
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Asundi, Vinod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/10119428
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Wehrman, Tom
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508.00
100.00%
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Best Local Similarity:
Query Match:
                                                                                                                                                                                         Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                          Score:
                                                                                                                                                                                                                                             Alignment Scores: Pred. No.:
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                                                                                                                            US-09-801-115B-2 (1-99) x US-09-918-995-19489 (1-512)
                                                                                                                                                                           Query Match:
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 19489
LENGTH: 512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR PILING DATE: 1999-01-20
PRIOR FILING DATE: 1999-01-20
                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (1) ... (512)
OTHER INFORMATION: n =
                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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202 GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                               142 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
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                21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
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                                                                                  MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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Matches:
Conservative:
Mismatches:
Indels:
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Matches:
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CURRENT APPLICATION NUMBER: US/10/319,763
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: 60/066,677
PRIOR FILING DATE: 1997-11-13
PRIOR PELICATION NUMBER: 60/069,957
PRIOR PELING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR FILING DATE: 1998-04-13
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR PILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR APPLICATION NUMBER: 60/099,273
PRIOR PILING DATE: 1998-09-04
PRIOR PILING DATE: 1998-09-04
PRIOR PILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
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SOFTWARE: Patent.pm
SEQ ID NO 53
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Publication No.
                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: 197..412
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
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APPLICANT: Duclert, Aymeric
APPLICANT: Bougueleret, Lydie
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: sig_peptide
LOCATION: 199..288
OTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 5.5999999046325
OTHER INFORMATION: seq IVSVLALIPETTT/LT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
NAME/KEY: misc_feature
LOCATION: 425..488
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 61.195
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: polyA site LOCATION: 489..500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: polyA_signal LOCATION: 464..469
                                                                                                                                                 OTHER INFORMATION: est
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Bougueleret, Lyd
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FEATURE:
NAME/KEY: misc feature
LOCATION: 425..488
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA516431
OTHER INFORMATION: est
                                                                                                                           NAME/KEY: misc_feature
LOCATION: 197..324
OTHER INFORMATION: homo:
OTHER INFORMATION: id ::
OTHER INFORMATION: est
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LOCATION: 61.195
OTHER INFORMATION: homology
OTHER INFORMATION: id:AA455042
OTHER_INFORMATION: est
               OTHER INFORMATION: homology
OTHER INFORMATION: id: W38899
OTHER INFORMATION: est
                                                                                                                                                                                                                                           NAME/KEY: miso_feature
LOCATION: 52..195
OTHER INFORMATION: homology
OTHER INFORMATION: id :W388!
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OTHER INFORMATION: homology
OTHER INFORMATION: id :W93646
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LOCATION: 58..195
OTHER INFORMATION: homology
OTHER INFORMATION: id :W93646
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LOCATION: 207..412
OTHER INFORMATION: homology
OTHER INFORMATION: id: W93646
OTHER INFORMATION: est
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OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION: id :AA45
OTHER INFORMATION: est
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NAME/KEY: misc_feature
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OTHER INFORMATION: homology
OTHER INFORMATION: id :AA51
OTHER INFORMATION: est
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature LOCATION: 90..195
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LOCATION: 197..412
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LOCATION: 197..412
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                                                                         OCATION: 443..477
EATURE:
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id :AA455042
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                                                                                                                                                 id :W38899
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                                                                                            US-10-242-535A-7940
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Best Local Similarity:
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                                           Sequence 7940, Application US/10242535A Publication No. US20040013663A1 GENERAL INFORMATION:
APPLICANT: ChondroGene Inc.
APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: 9..10,12
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LOCATION: 40..195
OTHER INFORMATION: homology
OTHER INFORMATION: id:W19506
OTHER INFORMATION: est
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LOCATION: 339..401
OTHER INFORMATION: homology
OTHER INFORMATION: id:w52820
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LOCATION: 71..195
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LOCATION: 425..469
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                                                                                                                                                            uPheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
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ORGANISM: Human
FEATURE:
NAME/KEY: misc feature
LOCATION: (176)...(176)
OTHER INFORMATION: n is
US-10-242-535A-7940
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CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR PILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR TILING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
SOPTWARE: PATENTIN PATENTING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
SOPTWARE: PATENTIN VERSION 3.2
SEQ ID NO 7940
LENGTH: 400
APPLICANT: CHEMOKINE LINE TITLE OF INVENTION: CHEMOKINE LINE TITLE OF INVENTION: STIMULATING ACTIVITIES; TITLE OF INVENTION: STIMULATING ACTIVITIES; FILE REFERENCE: 10776-003-999

CURRENT APPLICATION NUMBER: US/09/801,115; CURRENT FILING DATE: 2001-03-07

PRIOR APPLICATION NUMBER: PCT/CN00/00026
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Best Local Similarity:
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Patent No. US20020001828A1
GENERAL INFORMATION:
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APPLICANT:
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                                                                                                                                                             Ma, D.
Han, W.
Shang, Y.
Song, Q.
Di, C.
Huang, J.
Tang, J.
Chen, G.
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                                                                                                                         CHEMOKINE LIKE FACTOR (CKLF) WITH CHEMOTACTIC STIMULATING ACTIVITIES
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; PRIOR APPLICATION NUMBER: 99107284.7
; PRIOR FILING DATE: 1999-05-14
; NUMBER OF SEQ ID NOS: 8
; SEQ ID NO 3
; LENGTH: 459
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-801-115-3
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Pred. No.:
 421
                                                         361 TGTCTTGCCGACGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTAC 420
                                                                                                                 301
                                                                                                                                                                           241 ATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGCACTGATA 300
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                                                                                                                                 48 ProGluThrThrThrLeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCys 67
                         88
                                                                       68 CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr 87
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                GlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                                 CCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGC 360
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Search completed: January 31, 2004, 05:31:41 Job time: 275 secs

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-Q=/cgn2_1/USPTO_spool_p/US09801115/runat_30012004_113849_7051/app_query.fasta_1.263
-DB=Issued_patents_NA -QFMT=fastap -SUFFIX=p2n.rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DCCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15
-MODEL-CCAL -OUTPMT=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09801115_@CGN 1 1 103 @runat_30012004_113849_7051 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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-MODEL=frame+_p2n.model -DEV=xlp
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
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5: /cgn2_6/ptodata/2/ina/BTUS_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/backfiles1.seq:*
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Ygapop 10.0 , Ygapext
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US-09-599-360B-29
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US-09-599-360B-70
US-09-385-982-95
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US-09-251-645-11
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US-09-170-496D-179
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Sequence 98, Appl
Sequence 252, App
Sequence 29, Appl
Sequence 164, App
Sequence 53, Appl
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PP. 2	, Ař	Sequence 2, Appli Sequence 1, Appli Sequence 376, App Sequence 108, App	Sequence 7778, Ap Sequence 1, Appli Sequence 1, Appli Sequence 7562, Ap	Sequence 176, App Sequence 8, Appli Sequence 8, Appli Sequence 6, Appli	Sequence 1, Appli Sequence 2920, Ap Sequence 70, Appl Sequence 39, Appl	1, 9, 1, 14,	27 10 2, 6 2, 6 12,

ALIGNMENTS

RESULT 1 US-09-663-600A-147 ; Sequence 147, Appl ; Patent No. 6573068

Sequence 147, Application US/09663600A

INFORMATION:

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APPLICANT: Dumas Milne Edwards, Jean-Baptiste
APPLICANT: Duclert, Aymeric
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS
FILE REFERENCE: 31.US3.CIP
CURRENT APPLICATION UNMER: US/09/663,600A
CURRENT APPLICATION UNMER: 09/191,997
PRIOR FILING DATE: 1998-11-13
PRIOR APPLICATION NUMBER: 60/066,677
PRIOR FILING DATE: 1998-11-13
PRIOR APPLICATION NUMBER: 60/069,957
PRIOR FILING DATE: 1997-12-17
PRIOR FILING DATE: 1997-12-17
PRIOR FILING DATE: 1997-12-17
PRIOR FILING DATE: 1998-00-09
PRIOR APPLICATION NUMBER: 60/074,121
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: 60/081,563
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR APPLICATION NUMBER: 60/096,116
PRIOR FILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 229
SEQ ID NO 147
LENGTH: 413
TYPE: DNA
ORGANISM: Homo sapiens
PRATURE:
NAME/KEY: Sig_peptide
LOCATION: 46:.189
OTHER INFORMATION: Von Heijne matrix
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APPLICANT: ROSSEN et al.
TITLE OF INVENTION: 186 Human Secreted prote
FILE REFERENCE; PZ002P1
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER APPLICATION NUMBER: 60/040,333
EARLIER APPLICATION NUMBER: 60/038,621
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,626
EARLIER APPLICATION NUMBER: 60/040,626
EARLIER APPLICATION NUMBER: 60/040,334
EARLIER APPLICATION NUMBER: 60/040,334
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EARLIER APPLICATION NUMBER: 60/047,600
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EARLIER APPLICATION NUMBER: 60/047,615
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; LOCATION: 402..713
US-09-663-600A-147
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Query Match:
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Patent No.
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OTHER INFORMATION: seq VFMLIVSVLALIP/ET
NAME/KEY: polyA signal
LOCATION: 377..382
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ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,582
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,596
ER APPLICATION NUMBER: 60/047,612
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,612
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,632
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,601
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/043,580
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,580
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,314
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ER APPLICATION NUMBER: 60/043,315
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/048,974
ER FILING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/056,886
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,877 ER APPLICATION NUMBER: 60/043,674
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,669
ER FILLING DATE: 1997-04-11
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APPLICATION NUMBER: 60/056,637

1997-08-1997-08-

NUMBER: 60/056,882

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1997-08-

APPLICATION NUMBER: 60/056,903 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,888

APPLICATION NUMBER: 60/056,894 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,911

LING DATE:

1997-08-22

APPLICATION NUMBER: 60/056,879 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,880

LING DATE:

1997-08-22

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NUMBER: NUMBER:

60/056,872 60/056,662

1997-08-22

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TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P1
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: PCT/US98/04493
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER APPLICATION NUMBER: 60/040,333
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,333
EARLIER APPLICATION NUMBER: 60/040,333
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Best Local Similarity:
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US-09-149-476-252
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SR FILING DATE: 1997-08-22
SR APPLICATION NUMBER: 60/048,964
SR FILING DATE: 1997-06-06
SR APPLICATION NUMBER: 60/057,650
SR FILING DATE: 1997-09-05
SR FILING DATE: 1997-09-05
SR APPLICATION NUMBER: 60/056,884
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FILING DATE: 1997-09-05
APPLICATION NUMBER: 60/049,610
FILING DATE: 1997-06-13
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APPLICATION NUMBER: 60/056,887
FILING DATE: 1997-08-22
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APPLICATION NUMBER: 60/056,875
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,862
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APPLICATION NUMBER: 60/047,588

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LING DATE: LING DATE: APPLICATION NUMBER: 60/057,761 FILING DATE: 1997-08-22

APPLICATION NUMBER: 60/047,595

FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,599

FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,892

LING DATE:

1997-08-22

PPLICATION NUMBER: 60/056,845

1997-08-22

LING DATE:

FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,631

FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,910
APPLICATION NUMBER: 60/056,864

TLING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,874 PPLICATION NUMBER: 60/056,636

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E: 1997-08-22 N NUMBER: 60/056,881 E: 1997-08-22 N NUMBER: 60/056,909

FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,664
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,876

FILING DATE: APPLICATION I

NUMBER: 60/056,632

1997-04-11 1997-05-23

APPLICATION NUMBER: 60/043,670

APPLICATION NUMBER: 60/047,501

1997-04-11

LING DATE:

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FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,576 APPLICATION NUMBER: 60/043,578 FILING DATE: 1997-04-11 FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,614 APPLICATION NUMBER: 60/047,593 FILING DATE: 1997-05-23

LING DATE:

1997-05-23

FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,589 APPLICATION NUMBER: 60/047,594

LING DATE: 1997-05-23

APPLICATION NUMBER: 60/047,590

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ER PILLING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040, 163
ER FILLING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/047, 600
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047, 597
ER APPLICATION NUMBER: 60/047, 597
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047, 597
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047, 597
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047, 613
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047, 613
ER APPLICATION NUMBER: 60/047, 613
ER APPLICATION NUMBER: 60/047, 613
ER APPLICATION NUMBER: 60/047, 593
ER APPLICATION NUMBER: 60/043, 583
ER APPLICATION NUMBER: 60/043,

R FILING DATE:
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NUMBER: 60/047,501

1997-04-11 1997-05-23

APPLICATION NUMBER: 60/043,670

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DB:
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Best Local Similarity:
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards,
APPLICANT: Bougueleret, L.
                                                             Sequence 29, Application Patent No. 6548633
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APPLICATION NUMBER: 60/056,887
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,908
FILING DATE: 1997-08-22
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FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/057,650
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APPLICATION NUMBER: 60/056,875
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,862
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APPLICATION NUMBER: 60/061,060
FILING DATE: 1997-10-02
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APPLICATION NUMBER: 60/056,909
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APPLICATION NUMBER: 60/056,881
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APPLICATION NUMBER: 60/056,876
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FILING DATE: 1997-08-22
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Mismatches:
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Gaps:
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Percent Similarity:
Best Local Similarity:
Query Match:
; Sequence 164, Application US/0922013
patent No. 6506607
GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
TITLE OF INVENTION: METHODS AND CO.
TITLE OF INVENTION: OF PROSTATE CO.
FILE REFERENCE: 07334-074001
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NAME/KEY: sig_peptide
LOCATION: 144..287
COTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 4.10
OTHER INFORMATION: seq VFMLIVSVLALIP/ET
NAME/KEY: polyA signal
LOCATION: 457..462
NAME/KEY: polyA_site
LOCATION: 500..515
NAME/KEY: misc_feature
1000ation: 60
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PRIOR FILING DATE: 1998-12-22
PRIOR PPLICATION NUMBER: 60/141,032
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 123
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SEQ ID NO 29
LENGTH: 515
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TITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides FILE REFERENCE: GENSET 050CP3
CURRENT APPLICATION NUMBER: US/09/599,360B
CURRENT FILING DATE: 2000-06-21
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TYPE: DNA
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OTHER INFORMATION: n=a,
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GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, Jean-Baptiste

APPLICANT: Duclert, Aymeric

APPLICANT: Duclert, Lydie

TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS

FILE REFERENCE: 31.US3.CIP

CURRENT APPLICATION NUMBER: US/09/663,600A

CURRENT FILING DATE: 2000-09-15

PRIOR APPLICATION NUMBER: 09/191,997

PRIOR APPLICATION NUMBER: 09/066,677

PRIOR FILING DATE: 1998-11-13

PRIOR APPLICATION NUMBER: 60/066,957

PRIOR FILING DATE: 1997-11-13

PRIOR APPLICATION NUMBER: 60/069,957

PRIOR APPLICATION NUMBER: 60/074,121

PRIOR APPLICATION NUMBER: 60/074,121

PRIOR APPLICATION NUMBER: 60/074,121

PRIOR APPLICATION NUMBER: 60/074,121

PRIOR APPLICATION NUMBER: 60/074,121
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Best Local Similarity:
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(526)
OTHER INFORMATION: n = A,T,C o:
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PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/220,132
CURRENT FILING DATE: 1998-12-23
PRIOR APPLICATION NUMBER: US 60/079,303
PRIOR FILING DATE: 1998-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.:
              DR APPLICATION NUMBER: 60/074,121
DR FILLING DATE: 1998-02-09
DR APPLICATION NUMBER: 60/081,563
DR FILLING DATE: 1998-04-13
DR APPLICATION NUMBER: 60/096,116
DR FILLING DATE: 1998-08-10
APPLICATION NUMBER: 60/099,273
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o. 6573068
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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PRIOR FILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 229
SOFTWARE: Patent.pm
SEQ ID NO 53
SEQ ID NO 53
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OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 61..195
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 425..488
OTHER INFORMATION: id :AA429945
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 425..488
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 425..488
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LOCATION: 207.412
OTHER INFORMATION: homology
OTHER INFORMATION: id: W93646
OTHER INFORMATION: est
NAME/KEY: misc feature
LOCATION: 58.195
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NAME/KEY: misc feature
LOCATION: 425.-488
OTHER INFORMATION: homology
OTHER INFORMATION: id :AA455042
OTHER INFORMATION: est
                                                                                                    NAME/KEY: misc_feature
LOCATION: 197..412
OTHER INFORMATION: homology
OTHER INFORMATION: id:AA516431
OTHER INFORMATION: est
                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: 425..488
OTHER INFORMATION: homology
OTHER INFORMATION: id :W93646
OTHER INFORMATION: est
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OTHER INFORMATION: id
OTHER INFORMATION: est
                                        NAME/KEY: misc feature
LOCATION: 90..195
OTHER INFORMATION: homology
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LOCATION: 61..195
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NAME/KEY: misc_feature
LOCATION: 197..412
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INFORMATION: est

homology id :AA516431

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DB:
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Best Local Similarity:
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OTHER INFORMATION: id: W52820
OTHER INFORMATION: est
NAME/KEY: misc_feature
LOCATION: 339..401
OTHER INFORMATION: homology
OTHER INFORMATION: id: W52820
OTHER INFORMATION: est
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LOCATION: 197..338
OTHER INFORMATION: homology
OTHER INFORMATION: id:W52820
OTHER INFORMATION: est
NAME/KEY: misc_feature
LOCATION: 71..195
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LOCATION: 9..10,12
OTHER INFORMATION: n=a,
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LOCATION: 425.7469
OTHER INFORMATION: homology
OTHER INFORMATION: id :W52820
OTHER INFORMATION: est
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OTHER INFORMATION: id: w38899
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LOCATION: 443..477
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LOCATION: 197..324
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OTHER INFORMATION: id :W38899
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LOCATION: 52..]
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                                                   252
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                  60
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                                                                    eValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPh 60
eAlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLe
                                                   CGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTT
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id:W19506
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Matches:
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FILE REFERENCE: GENSET 050CP3
CURRENT APPLICATION NUMBER: US/09/599,360B
CURRENT FILING DATE: 2000-06-21
PRIOR APPLICATION NUMBER: 60/113,686
PRIOR PILING DATE: 1998-12-22
PRIOR APPLICATION NUMBER: 60/141,032
PRIOR FILING DATE: 1999-06-25
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR APPLICATION NUMBER: 09/469,099
PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 123
SOFTWARE: PATENT. pm
SEQ ID NO 70
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Best Local Similarity:
Query Match:
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NAME/KEY: SIG_Deptide
LOCATION: 140.442
OTHER INFORMATION: Von Heijne matrix
OTHER INFORMATION: score 4.10
OTHER INFORMATION: seq VFMLIVSVLALIP/ET
NAME/KEY: polyA signal
LOCATION: 630..635
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Jobert, S.
TITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bougueleret, APPLICANT: Jobert, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dumas Milne Edwards, J.B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: polyA_site LOCATION: 655..669
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NAME/KEY: CDS
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                               28
                                                                                                                                                                                                                                              21 ValLysMetLeuArg-----
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<u>ÁTTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATA</u>
                      IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle
                                                                   CTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCTTGAT
                                                                                                                                         GCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTTCATA 319
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65.13%
65.13%
92.81%
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Matches:
Conservative:
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Indels:
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NUMBER OF SEQ ID NOS: 544
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 95
LENGTH: 392
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(392)
OTHER INFORMATION: n = A,T,C or G
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US-09-251-645-11
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DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOVEL HUMAN GENES AND GE
TITLE OF INVENTION: PRODUCTS: II
FILE REFERENCE: CCDNA-260XX
CURRENT APPLICATION NUMBER: US/09/385,982
CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1999-08-08
EARLIER FILING DATE: 1998-08-31
NUMBER: 60 EGO 11 NOSC. 544
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              Sequence 11, Application US/09251645 Patent No. 6281413
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity:
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INFORMATION:
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                                                                                                                      ProTyrGlnLysLysProValHisGluLysLys
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RESULT 10
US-09-116-498-11
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DB:
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Best Local Similarity:
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APPLICANT: Morgan, Michael K.
APPLICANT: Anderson, Arne R.
APPLICANT: Hart, Hope
APPLICANT: Warren, Gregory W.
APPLICANT: Dunn, Martha
APPLICANT: Chen, Jeng S.
                                                                                                    Sequence 11, Application US/09116498
Patent No. 6251582
GENERAL INFORMATION:
APPLICANT: Littman, Dan R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NOVEL INSECTICIDAL TOXINS FROM PHOTORHABDUS TITLE OF INVENTION: AND NUCLEIC ACID SEQUENCES CODING THEREFOR FILE REFERENCE: CGC1963/A CURRENT APPLICATION NUMBER: US/09/251,645
CURRENT FILING DATE: 1999-02-17
NUMBER OF SEQ ID NOS: 22
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NAME/KEY: CDS
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NAME/KEY: CDS
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NAME/KEY: CDS
LOCATION: (15171)..(18035)
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Unitimaz, Derya
Ramani, Vineet N.K.
ROVEL ALTERNATIVE G-COUPLED RECEPTORS
INVENTION: NOVEL ALTERNATIVE G-COUPLED RECEPTORS
ASSOCIATED WITH RETROVIRAL ENTRY INTO
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Matches:
Conservative:
Mismatches:
Indels:
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ORGANISM: pigtail macaque;
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-116-498-11
             RESULT 11
US-09-170-496D-179
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Best Local Similarity:
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Sequence 179, Application US/09170496D
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REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 104*
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson,
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ATTORNEY/AGENT INFORMATION:
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                                                               CAGCAGTCAGGAAAGCACAACAAAAAG 705
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                                                                                                                                                           CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr 87
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                                                                                          GlnLysLysProValHisGluLysLys
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STRANDEDNESS: double
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US-09-116-498-7
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CURRENT FILING DATE: 1998-10-13
NUMBER OF SEQ ID NOS: 294
SOFTWARE: PatentIn version 3.1
SEQ ID NO 179
LENGTH: 1083
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GENERAL INFORMATION:
APPLICANT: Behan, Dominic P.
                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/09116498 Patent No. 6251582
GENERAL INFORMATION:
APPLICANT: Littman, Dan R.
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ZIP: 07601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson,
STREET: 411 Hackensack Ave,
                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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                                                                                               CITY: Hackensack
STATE: New Jersey
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                                                                               COUNTRY:
                                                                                                                                                                                                                                                     INVENTION: NOVEL ALTERNATIVE G-COUPLED RECEPTORS
ASSOCIATED WITH RETROVIRAL ENTRY INTO CELLS, METHODS
IDENTIFYING THE SAME, AND DIAGNOSTIC AND THERAPEUTIC
                                                                                                                                                                                                                                                                                                    Unutmaz, Dergantmani, Vineet N.K.
                                                                                                                                                                                                                                                                                                                                                          Deng,
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Matches:
Conservative:
Mismatches:
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Best Local Similarity:
Query Match:
DB:
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US-09-170-496D-27
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CURRENT APPLICATION NUMBER: US/09/170,496D
CURRENT FILING DATE: 1998-10-13
NUMBER OF SEQ ID NOS: 294
SOFTWARE: PatentIn version 3.1
SEQ ID NO 27
LENGTH: 1083
                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                    Sequence 27, Application US/09170496D Patent No. 6555339
                                                                                                       APPLICANT: Behan, Dominic P.
APPLICANT: Chalmers, Derek T.
APPLICANT: Liaw, Chen W.
TITLE OF INVENTION: No. 6555339-Endogenous,
TITLE OF INVENTION: Receptors
FILE REFERENCE: AREN-0040
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SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA
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REFERENCE/DOCKET NUMBER: 10-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
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APPLICATION NUMBER: US/09/116,498
FILING DATE: 16-Jul-1998
CLASSIFICATION: JUNKnown>
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SOFTWARE: Patentin Release #1.0, Version #1.30
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71.00
39.33%
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DB:
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US-09-016-434-1079
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                    TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO:
                                                                     CLASSIFICATION:
ATTORNEY AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION UNMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
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APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
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     SEQUENCE CHARACTERISTICS:
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                                                        TELEPHONE:
                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: PALO ALTO
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                                                        (650) 855-0555
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Matches:
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US-09-016-434-1079
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Best Local Similarity:
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                 CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: Meuth, Donna M
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 0187
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
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LIBRARY: GENBA
                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                              APPLICATION NUMBER: FILING DATE: 14-MAN CLASSIFICATION: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ITLE OF INVENTION: ANTI-erbB-2 ANTIBODIES, COMBINATIONS
ITLE OF INVENTION: THEREOF, AND THERAPEUTIC AND DIAGNOSTIC USES THEREOF
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                                                                                                                                                                                                                                                                                                          COUNTRY: United States ZIP: 22313-1404
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BIRD, Robert E
(703) 836-2021
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LENGTH: 720 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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522 ACCATTCTCAGGATGAATCCATCCAATCCATTCCAG 487
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                                                                                    ValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLys--
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-DB=N Geneseq 19Jun03 -OFMT=fastap -SUFFIX=p2n.rng -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=biosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR SCORE-pct -THR MAX=100 -THR MIN-0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09801115 @CGN 1 1 490 @runat 30012004 113847 7015 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQÜERY -NGE_SCORES=0 -WAIT -DSPELOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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SUMMARIES

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Breast cancer rela	Breast cancer rela	cancer	744 used	nsmem	Human secreted exp	Human secreted pro	Human PRO772 EST p	no.			\sim	UCK-3 nucleotide s	cDNA encoding nove		UCK-4 nucleotide s	Human colon cancer	Human secreted pro	cDNA sequence #275	cDNA encoding nove	n polynuc	cDNA encoding huma	n secreted	cDNA encoding huma	Human PRO772 nucle	Human PRO772 (UNQ4		n transmembra	nucleotic	secreted	protein	-	ጠ .			n EST-derived		D.	Secreted		secreted		secreted	sequenc	uman secreted	Description

ALIGNMENTS

AAX97873 standard; cDNA; 413 BP.

AAX97873;

23-SEP-1999 (first entry)

Human secreted protein encoding cDNA #61.

AAXS9783
ID AAXX9783
ID AAXX
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AC AAXX
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XX BECL
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XX WO99 Secreted protein; human; cytostatic; diagnostic; gene therapy; chromosome mapping; secretion thrombotic; osteopathic; forensic;

Homo sapiens

WO9925825-A2

Pred. No. is the number of results score greater than or equal to the

predicted by chance to have a score of the result being printed,

27-MAY-1999

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RESULT 2
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13-APR-1998;
10-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used to express secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures. Uses also include design of expression vectors and secretion vectors.
  Human secreted protein; hyperproliferative disorder; autoimmune
                         cDNA sequence #397 encoding novel human secreted protein.
                                                  08-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 other;
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0
0
0
   disorder;
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                                                                                                                                                                                                285
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infectious disorder; allergic condition; neurodegenerative disorder; liver fibrosis; coagulation disorder; gene therapy; antimicrobial; tumour; cancer; hepatotronic; immunocurations.
    cancer; hepatotropic; immunosuppressive; antirheumatic; gene;
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Homo sapiens

WO200177289-A2

29-MAR-2001; 2001WO-US10232

06-APR-2000; 2000US-195605P

GENETICS INST

Jacobs K, n Merberg D, Clark HF, Treacy M, Fechtel K, мссоу лм, Lavallie ER
, Agostino M
, Howes SH, ER, COLLING 10 MJ, Bowman MR, Spaulus 3H, Resnick RJ, Gulukota Collins-Racie LA, Eva...
Dowman MR, Spaulding V Wong GG;

WPI; 2002-179322/23.

Six hundred and twenty three polynucleotides derived from a variety of human tissue sources which encode secreted proteins, useful for treating immune deficiencies and disorders such as autoimmune disorder disorders

Claim 1; Page 296-297; 393pp; English.

immunodeficiency (SCID)), autoimmune disorders (e.g. multiple sclerosis), blood disorders (e.g. thrombocytopaenia), inflammatory disorders (e.g. arthrittis), infectious disorders (e.g. hepatitis), allergic conditions (e.g. asthma), neurodegenerative disorders (e.g. Alzheimer's disease), liver fibrosis, coagulation disorders (e.g. Alzheimer's disease), liver fibrosis, coagulation disorders (e.g. derived from a variety of human tissues. The invention also provides a method for producing proteins from these polynucleotide sequences. The proteins are useful for identifying compounds that modulate their activity and production. The sequences of the invention are useful for treating diseases such as hyperproliferative disorders (e.g. cancer), immune deficiency disorders (e.g. severe combined immunodeficiency (SCID)), autoimmune disorders (e.g. multiple which encode human secreted derived from a variation of haemophilia), and tumours. The polynucleotide sequences of the invention are also useful in gene therapy. ABK35610-ABK36232 rejthe cDNA sequences of the invention that encode for novel human The present invention relates to the isolation of novel cDNA sequences secreted proteins. proteins. The cDNA sequences have represent

Sequence 439 B₽; 117 Þ 96 Ç 104 ູດ 122 Η. 0 other;

GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT	ACCAGAAACCACAACATTG/	TTGGCACTGATA	1 GTATCTGTG	b 201	Дb
ValSerValLeuAlaLeuIleProGluThrThrLeuThrValGlyGlyGlyValPhe	eProGluThrThrThrLeuf	LeuAlaLeuIl 	ValSerVal	y 41	Ş
GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC	TATTATCAACTCACTGGTA	CTGCGGCTGGA	1 GTGAAGATG	1	DЬ
21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle	pIleIleAsnSerLeuVal:	LeuArgLeuAsı	1 ValLysMet		δ
ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC	AATAAAACATCGCCCCTTC	STGCAGCCGAA	1 ATGGATAAC	b 81	뮍
MetAspAsnValGlnProLysI1eLysHisArgProPheCysPheSerValLysGlyHis :	sIleLysHisArgProPhe(ValGlnProLys	MetAspAsn	Y 1	8
	(1-439)	x ABK36006	5B-2 (1-99)	US-09-801-115B-2 (1-99) x ABK36006 (1-439)	c
0	Gaps:	24		DB:	b
0	Indels:	100.00%		Query Match:	o
0	Mismatches:	100.00%	imilarity:	Best Local Similarity:	B
0	Conservative:	100.00%	larity:	Percent Similarity:	שי
99	Matches:	508.00		Score:	တ
439	Length:	5.93e-58		Pred. No.:	שי
			ores:	Alignment Scores:	Α

5

AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu

260 60 200 40 140 20

밁 5 В

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02-OCT-1997;
07-MAR-1997;
07-MAR-1997;
07-MAR-1997;
07-MAR-1997;
07-MAR-1997;
07-MAR-1997;
 11-APR-1997
11-APR-1997
11-APR-1997
11-APR-1997
11-APR-1997
23-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteocporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-1997;
07-MAR-1997;
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11-APR-1997;
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11-APR-1997;
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11-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
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 97US-0040336.
97US-0040331.
97US-0043313.
97US-0043313.
97US-0043569.
97US-0043576.
97US-0043576.
97US-0043578.
97US-0043579.
97US-0043579.
97US-0043571.
97US-0043571.
97US-0043571.
97US-0043573.
97US-004759.
97US-004759.
97US-004759.
97US-004759.
97US-004758.
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97US-0038621.
97US-0040161.
97US-0040162.
97US-0040162.
97US-0040333.
97US-00403334.
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    23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
23-MAY-1997
16-JUN-1997
16-JUL-1997
16-JUL-1997
22-AUG-1997
23-AUG-1997
23-AUG
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23-MAY-1997
23-MAY-1997
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23-MAY-1997;
                                                                               Bednarik I
Feng P, I
Kyaw H, I
Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAY-1997;
23-MAY-1997;
23-MAY-1997;
23-MAY-1997;
New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological
                                        WPI; 1998-506364/43.
P-PSDB; AAW74818.
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23-MAY-1997
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                                                                                                                                                (HUMA-) HUMAN
                                                                            k DP, Brewer LA, C
Ferrie AM, Fische
Lafleur DW, Li Y,
M, Shi Y, Soppet C
                                                                                                                                                GENOME
                                                                                                                                                                                                                                                                                                                                     97US-0056874.
97US-0056877.
97US-0056875.
97US-0056877.
97US-0056877.
97US-0056878.
97US-0056880.
97US-0056880.
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97US-0056880.
97US-0056886.
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97US-0051926
97US-0055874
97US-0056530
97US-0056630
97US-0056631
97US-0056631
97US-0056636
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97US-0047595.
97US-0047597.
97US-0047599.
97US-0047600.
97US-0047601.
97US-0047613.
97US-004761.3.
97US-004761.3.
97US-004761.3.
97US-004761.3.
97US-004763.3.
97US-0048974.
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97US-0056909.
97US-0056910.
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97US-0056893.
97US-0056894.
97US-0056903.
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97US-0056862.
97US-0056864.
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97US-0056664
                                                                           r LA, Carter KC, Duan R, Ebner R, E
Fischer CL, Florence KA, Greene JM,
1, Li Y, Moore PA, Ni J, Olsen HS,
Sommet DR, Young PE, Yu GL, Zeng Z;
                                                                                                                                                SCI INC
                                                                         CL,
Moore PA, N.
Young PE,
                                                                                   Ebner R, Knurca
Greene JM, Hu JS;
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RESULT 4
ARS73E ID AB87
ID AB87
XX ARS7
XX ARS7
XX Huma
XX Hum
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a nucleic acid molecule designated Gene 88 from the human cDNA clone HADAV32 (deposited as clone ATCC 97901 and ATCC 209047) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human clone fusion proteins by linking to the gene to a human from the fused protein as compared to the human protein only. The invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAV74731-W75026) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypeptides in a golynucleotides. Specific uses are described for each of the 186 polynucleotides, based on which tissues they are most highly expressed in
                                                                                         Human; ss; gene; secreted protein; autoimmune disease; chemotaxis; rheumatoid arthritis; hyperproliferative disorder; breast neoplasm; liver neoplasm cardiovascular disorder; cardiac arrest; skin aging; cerebrovascular disorder; cerebral ischaemia; angiogenesis; sunburn; nervous system disorders; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; tissue regeneration; epithelial cell proliferation; organ transplantation; food additive;
                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 504
                                                                                                                                                                                                                                                                                            Human cDNA
                                                                                                                                                                                                                                                                                                                                                                                        ABS73585
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                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; cDNA; 504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
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                                                                        nutritional
                                                                                                                                                                                                                                                                                         for novel secreted protein gene
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508.00
100.00%
100.00%
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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    23-MAY 1997
23-MAY
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23-MAY-1997
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9718-040333P.
9718-04033P.
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9718-0433112P.
9718-043315P.
9718-043315P.
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9718-047802P.
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9718-047882P.
9718-047882P.
9718-047882P.
9718-047883P.
9718-047883P.
9718-047883P.
9718-047832P.
9718-047832P.
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(HUMA-) HUMAN GENOME SCI INC

Endress GA, Duan R, Hu Moore PA, MS, Shi Y, Rosen CA, 4 Yu G, Ni J, Feng J, Florence KA, C ni Y, Lafleur DW, Fischer CL, & Fengr, KA, Olsen Ho, n HS, EL. V. Zeng Z, Soppet DP, Carter KC,
Young PE, Greene JM,
n HS, Ebner R, Brewer
Y, Zeng Z, Kyaw H; Ę, Ferrie AM:

P-PSDB; ABG95267 WPI; 2002-634796/68.

New isolated human secreted protein for diagnosing, preventing, treating or ameliorating medical conditions and used as a food additive preservative

Example 1; SEQ ID ĕ 98; 129pp; English

carbohydrate, vitamins, minerals, colactors and control of the present sequence represents a cDNA derived from a components. The present sequence represents a cDNA derived from a gene encoding one of the novel human secreted proteins of the invention. Note: This sequence did not form part of the printed specification, but was obtained in electronic format directly from USPTO at angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, minerals, cofactors and other nutritional The invention relates to an isolated protein that is one of 186 human secreted proteins, given in the specification, encoded by one of 309 cDNA sequences also given in the specification. The protein is used in a pharmaceutical composition used to prevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia,

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97US-0040333.

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97US-0040626.

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97US-0043313.
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97US-0047613
97US-0056631
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97US-0056909
97US-0056910
97US-0057650
97US-0057650
97US-0057650
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(HUMA-) HUMAN GENOME SCI INC

Feng P, E Kyaw H, I Ruben SM, Bednarik DP, Ferrie AM, Fisch Lafleur DW, Li Y 1, Shi Y, Soppet Brewer IA, Carter KC, Duan R, Ebner R, ie AM, Fischer CI, Florence KA, Greene JM eur DW, Li Y, Moore PA, Ni J, Olsen HS, i Y, Soppet DR, Young PE, Yu GL, Zeng Z; ı Y, DR, JM, LL Rosen C Endress GA; M, Hu JS; Rosen CA;

WPI; 1998-506364/43. P-PSDB; AAW74961.

New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders

Claim 1; Page 472; 721pp; English.

This sequence represents a nucleic acid molecule designated Gene 88 from CC the human cDNA clone HALAV32 (deposited as clone ATCC 97897 and ATCC 209043) which encodes a secreted human protein. The gene can be used to CC generate fusion proteins by linking to the gene to a human community of the fused protein as compared to the human protein only.

CC the fused protein as compared to the human protein only.

CC the invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAW7471-W75026)

CC which are useful for preventing, treating or ameliorating medical CC conditions e.g. by protein or gene therapy. Also, pathological CC conditions can be diagnosed by determining the presence of mutations in CC polypeptides in a sample or by determining the presence of mutations in CC polynucleotides. Specific uses are described for each of the 186 CC polynucleotides, based on which tissues they are most highly expressed in CC (see AAV59511 for described uses). Sequence 506 BP; 148 A; 103 C; 121 G; 132 T; 2 mutations in ch of the 186 y expressed in

DB:	Query Match:	Best Local Similarity:	Percent Similarity:	Score:	Pred. No.:	Alignment Scores:
19	100.00%	100.00%	100.00%	508.00	7.19e-58	
Gaps:	Indels:	Mismatches:	Conservative:	Matches:	Length:	
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US-09-801-115B-2 (1-99) x AAV59746 (1-506)

82	Дb	8	Ф	Ş	В	Q
61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80	237 GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT 2	41 ValSerValLeuAlaLeuIleProGluThrThrThrThrTeuThrValGlyGlyValPhe 60	177 GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC 2	21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40	117 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC 1	1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis 20
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RESULT 6
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97US-040336P
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        23-MAY-1997
22-AUG-1997
23-AUG-1997
23-AUG
                                            WPI; 2002-634796/68.
P-PSDB; ABG95421.
                                                                                                      Ruben SM, Rosen CA, Fischer CL, Endress GA, Yu G, Ni J, Feng P, Duan R, Hu J, Florence KA, Olsen Moore PA, Shi Y, Lafleur DW, Li
    New isolated human secreted protein for diagnosing, preventing,
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97US-0585P85P.
97US-061060P.
98WO-US04493.
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                                                                                              Feng r,
KA, Olsen F
                                                                                                      Soppet DP,
, Young PE,
en HS, Ebner
i Y, Zeng Z,
                                                                                                        Carter KC,
Greene JM,
C R, Brewer
Kyaw H;
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RESULT 7
AAA87730
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AC AAA8
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia,
Human; secreted protein; forensic procedure; gene therapy; chromosome mapping; cancer; autoimmune disease; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          treating or ameliorating medical conditions and used as a food additive or preservative {\ \ }
                                                                                                      28-NOV-2000
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                                                            Human secreted protein encoding cDNA SEQ ID #29.
                                                                                                                                                                               AAA87730 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a pharmaceutical composition used to prevent, treat or ameliorate a dical condition in e.g. humans, mice, rabbits, goats, horses, cats, gs, chickens or sheep. Disorders which are diagnosed or treated include
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to an isolated protein that is one of 186 human reted proteins, given in the specification, encoded by one of CDNA sequences also given in the specification. The protein is us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAAGGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
                                                                                                                                                                                                                                                                                                                                                                                                                              GTATCTGTGTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
                                                                                                                                                                                                                                                                                                     PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
                                                                                                                                                                                                                                                                                                                                               GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG
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                                                                                                                                                                                                                                                                                               Score:
                                                                                                                                                                                                                                                                                                                                Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents human cDNA encoding a secreted protein. The invention relates to sequences AAA87725-A87774 which encode human secreted proteins AAB25763-B25812. The proteins include signal peptides. Included in the invention are a host cell containing one of the cDNA sequences, and a purified antibody capable of binding to one of the secreted proteins. Also contained in the invention are methods for storing the sequence data on a computer system, and a method for identifying features of the cDNA sequences using a computer programme. The cDNAs are useful for expressing secreted proteins or fragments to obtain antibodies capable of specifically binding to the secreted proteins. The cDNAs may also be useful in diagnostic, forensic, gene therapy and chromosome mapping procedures and may be used to design expression vectors and secretion vectors. The proteins of the invention expression vectors and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                    may be used to treat diseases including cancer, autoimmune diseases, cardiovascular disorders, cystic fibrosis, hypothyroidism, immunological disorders, amyloidosis, brain disorders, skeletal muscle disorders, eye disorders, obesity, mitochondriccytopathies, diabetes, atherosclerosis, neurodegenerative disorders, graft rejection, Alzheimer's disease, dementia, hyperlipidaemia, septic shock and impotence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cystic fibrosis; hypothyroidism; immunological disorder; amyloidosis; brain disorder; skeletal muscle disorder; eye disorder; obesity; mitochondriocytopathy; diabetes; atherosclerosis; Alzheimer's disease; neurodegenerative disorder; graft rejection; dementia; hyperlipidaemia;
                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bougueleret L,
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25-JUN-1999;
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ValSerValLeuAlaLeuIleProGluThrThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                     ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                    ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
                                                                                                                                      MetAspAsnValG1nProLys11eLysHisArgProPheCysPheSerValLysG1yHis
                                               GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
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99US-0141032
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to 49 Secreted proteins and the cDNAs encoding them. The protein and nucleic acids may be used in the prevention, treatment and diagnosis of diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            49 Secreted proteins and the nucleic acids encoding them, us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 515
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21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inappropriate protein expression.
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                                                                                                                                                                             Alignment Scores:
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                                                                                                                                                                                                                      This sequence represents an UCK-1 cDNA sequence encoding a chemotaxis factor polypeptide. The UCK-1 protein exhibits immunocyte chemotaxis activity and a haemopoiesis stimulating effect. The invention relates UCK proteins, their encoding nucleotide sequences and antibodies and antagonists against the proteins. The nucleotide and protein sequences are useful for the preparation of a composition for the diagnosis and treatment of diseases associated with abnormal immunocyte function and low haemopoiesis function caused by radiotherapy and chemotherapy used treat tumours and other diseases.
                                                                                                                                                                                                    Sequence 534 BP;
                                                                                                                                                                                                                                                                                                                                      Example 4; Fig 1; 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                           Chemotaxic factor useful for treatment and diagnosis of immunocyte disorders - has immunocyte chemotaxic stimulating factor
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GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
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17-JUL-2000;
03-AUG-2000;
15-SEP-2000;
                                                                                                                                                                                                                                                       proteins from a variety of organisms, including human, dog, cat, hor cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags of from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a feat of the present sequence is a feat of the present sequence.
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                                                                                                                                                                                                                                                                                                                                                                                The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated polypeptide for treatment antibodies and research use -
                                                                                                                                                                                          Sequence 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 451; 1275pp; English.
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                                                                     The present invention provides the protein and coding sequences of proteins from a variety of organisms, including human, dog, cat, ho cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JAN-2000; 2000US-0491404
17-JUL-2000; 2000US-0617746
03-AUG-2000; 2000US-0631451
15-SEP-2000; 2000US-0663870
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                  AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P
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03-NOV-1999;
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     N and P may be used to treat disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of p by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention.

N.B. Pages 666 to 682 and page 7053 of the missing at time of publication, meaning no SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                         Human; secreted protein, arthritis; Crohn's disease; sepsia; shock; ischaemia reperfusion injury; haematopoiesis; cancer; neuropathy; transgenic animal; Alzheimer's disease; Parkinson's disease; burn; amyotrophic lateral sclerosis; platelet disorder; thrombocytopenia;
                                                                                                                                                                                                                                                                                            18-DEC-2001
                                                                                                                                                                                                                                                                                                                        AAS44932
                                                                                                                                                                                                                                                                                                                                                  AAS44932 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 558 BP;
                                                                                                                                                                                                                                                                 encoding novel human secretory protein,
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ulcer; osteoporosis; bone degenerative disorder; periodontal disease; gut protection; lung; liver fibrosis; immune deficiency; infection; severe combined immunodeficiency; SCID; autoimmune disorder; allergy; multiple sclerosis; rheumatoid arthritis; diabetes mellitus; asthma;

fertility; analgesic; pain; antigen;

sapiens.

05-MAR-2001; 2001WO-US04942.

13-SEP-2001.

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В Ś В Š

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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                 CC ischaemia-reperfusion injury, shock, sepsis, immune responses, and is convolved in increasing haematopoiesis, stem cell survival, bone growth can remodeling. (I), (II) and modulators of (II) are useful for creating transgenic animals useful for studying the in vivo activities of the prohylaxis or treatment of one or more cancers. (II) is also useful for the polypeptide as well as for studying modulators of the polypeptides. (I) induces the proliferation of neural cells and regeneration of nerve can brain tissue and is useful for the treatment of central and corriberal nervous system diseases and neuropathies, such as Alzheimer's, partitions, the injury in various tissue and anyotrophic lateral cell sectivity, regulation of haematopoiesis and is useful for treating myeloid cor lymphoid cell disorders, platelet disorders such as thrombocytopenia cell sorders, or periodontal disease. Furthermore, (I) is also useful for treating cell disorders platelet disorders such as thrombocytopenia cell disorders, platelet disorders such as thrombocytopenia cell disorders, or periodontal disease. Furthermore, (I) is also useful for gut protection or regeneration and treatment of lung or liver fibrosis, ceperfusion injury in various tissues, various immune deficiencies and cell disorders including severe combined immunodeficiency (SCID), bacterial or fungal infections, autoimmune disorders e.g. multiple sclerosis, ce fungal infections, autoimmune disorders e.g. multiple sclerosis, ce faitatity, metabolism, catabolism, anabolism, storage or elimination of faitation, injects biorhythms or circadian cycles of rhythms, ce faitatist, mysters or witaning, minerals, provides and conditions, catabolism, anabolism, storage or elimination of content provides of catabolism.
                                                                                                                             US-09-801-115B-2 (1-99) x AAS44932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel polypeptides and nucleic acids obtained from prepared from various human tissues, for diagnosis cancer, neurological, inflammatory, and autoimmune
                                                                                                                                                                                                                                                                                                                                                                    activity and can act as an antigen in a vaccine composition to raise an immune response. AAS44920-AAS45295 represent novel human secreted prote:
                                                                                                                                                                                                                                                                                                                                                                                                      dietary fat, lipid, protein, carbohydrate, vitamins, minerals, provides analgesic effects or other pain reducing effects, immunoglobulin like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUL-2000;
19-SEP-2000;
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19-MAY-2000;
                                                                                                                                                                                                                                                                                                                 Sequence 655
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                                                                                                                                                                                                                                                              No.
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inflammatory conditions such as arthritis, nephritis, Crohn's disease,
ischaemia-reperfusion injury, shock, sepsis, immune responses, and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HYSE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention relates to novel isolated human secreted polypeptides (I)
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ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                        ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID No 13; 107pp; English.
                                                                             MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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; 2000US-0574454.
; 2000US-0596193.
; 2000US-0616847.
; 2000US-0665363.
; 2000US-0693267.
                                                                                                                                                                                                                                                                                                                   BP;
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ng J, Chen
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s and treatment o
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25-AUG-1998;
09-SEP-1998;
29-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytokine production; cell proliferation; cell differentiation; immune deficiency; infectious disease; autoimmune disorder; asthma; multiple sclerosis; systemic lupus erythmatosus; rheumatoid arthritis; allergic reaction; osteoporosis; osteoarthritis; periodontal disease; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; liver fibrosis; lung fibrosis; reperfusion injury systemic cytokine damage; tissue differentiation; contraceptive; stroke coagulation disorder; myocardial infarction; inflammatory condition; septic, shock; sepsis; ischaemia; reperfusion injury; arthritis; tumour;
This sequence encodes a human protein of the invention, which has hydrophobic domains. The DNA sequences can be used as a probe or as a genetic marker. The protein can also be used as a marker, and to identify potential genetic disorders. The DNA and protein can also be used as nutritional sources or supplements. The protein can also be used as proliferation, cell differentiation activities and induces production of other cytokines in certain cell populations. The protein also exhibits immune stimulating or immune suppressing activity. It can be used in the treatment of various immune deficiencies and disorders, and to treat
                                                                                                                                                                                                          Novel human proteins having hydrophobic domains useful for treating osteoporosis, Alzheimer's disease, Parkinson's disease, asthma, multiple sclerosis, rheumatoid arthritis, cancer, anaemia, and stroi
                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                 P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                     (SAGA )
(PROT-)
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07-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nephritis; therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUN-2000
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PROTEGENE INC.
                                                                                                                                                                        Page 217-218; 351pp;
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98JP-0254736.
98JP-0275505.
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                                                                                                                                                                                                                                                                                                                                                                                     CHEM RES CENT
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98JP-0224105
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conditions such as asthma, and in immune suppression after organ transplantation. The protein is useful in regulation of haematopoiesis and consequently in the treatment of myeloid or lymphoid cell deficiencies. It is also used in compositions for tissue growth or regeneration. The protein is also used in the treatment of osteoporosis or osteoarthritis and in the treatment of periodontal disease and other tooth repair processes. The protein is used in the treatment of nervous system disorders such as Alzheimer's disease, Parkinson's disease, and huntington's disease. They are useful for protection or regeneration and treatment of lung or liver fibrosis, repertusion injury in various tissues, and conditions resulting from systemic cytokine damage. They are also used for promoting or inhibiting tissue differentiation. They are also used as contraceptives since they exhibit activin or inhibin related activities and as a fertility inducing therapeutic. They are used for treatment of sorders and in treatment and prevention of infarction or stroke. They also acts as receptors, receptor ligands or inhibitors resulting from coagulation activities e.g. myocardial interactions. They are used to result in from a completions activities e.g. myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infectious diseases caused by viral, bacterial, fungal or other infections. The protein is also used for treating autoimmune dissuch as multiple sclerosis, systemic lupus erythematosus, and riarthritis. It is also useful in the treatment of allergic react.
reperfusion injury, arthritis,
                                                                 inflammatory conditions such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in the treatment of allergic reactions and
     such as septic shock, sepsis, ischaemia and nephritis. They can be used to
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d rheumatoid
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Sequence 297 BP; 81 A; 89 <u>ე</u> 72 <u>ი</u> 76 Η, 0 other;

No

Length: Matches:

98

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Best Local Similarity:
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                                       121
           61
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AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                        GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
                                                         ValSerValLeuAlaLeuIleProGluThrThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                               GTGAAGATGCTGCGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                                                                    MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
                                                                                                                                                    ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
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100.00%
98.99%
99.21%
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81 PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu

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181

GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG

99

240

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241

AAA15929 standard; cDNA; 467 ВP

12-JUN-2000 (first entry)

AAA15929

Human protein clone HP10357 full length coding sequence.

RESULT 15
AAA15929
ID AAA15
XX
AC AAA15
XX
TI 12-JU
TI 12-JU
XX
TI Human
XX
Human
KW Human
KW immun
KW immun
KW multck
KW multck
KW aller Human procein; hydrophobat volument, cell differentiation; cytokine production; cell proliferation; cell differentiation; immune deficiency; infectious disease; autoimmune disorder; asthma; multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis; multiple sclerosis; osteoportesis; osteoportesis; periodontal disease; Human protein; hydrophobic domain; nutritional source; haematopoiesis;

> nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; liver fibrosis; lung fibrosis; reperfusion injury systemic cytokine damage; tissue differentiation; contraceptive; stroke coagulation disorder; myocardial infarction; inflammatory condition; septic shock; sepsis; ischaemia; reperfusion injury; arthritis; tumour; is; reperfusion injury; contraceptive; stroke; injury;

sapiens.

WO200005367-A2

03-FEB-2000

22-JUL-1999; 99WO-JP03929

07-AUG-1998; 25-AUG-1998; 09-SEP-1998; 29-SEP-1998; 98JP-0238116 98JP-0254736 98JP-0275505 98JP-0208820 98JP-0224105

(PROT-) (SAGA) SAGAMI CHEM RES CENT PROTEGENE INC

2000-182694/16.

Novel human proteins having hydrophobic domains useful for treating osteoporosis, Alzheimer's disease, Parkinson's disease, asthma, multiple sclerosis, rheumatoid arthritis, cancer, anaemia, and stroi and stroke

Claim 4; Page 228; 351pp; English

Completive entrodes a number protein of the Environment of Andrews and the Sequences can be used as a probe or as a componential genetic disorders. The DNA and protein can also be used as a marker, and to identify potential genetic disorders. The DNA and protein can also be used as a multiple sources or supplements. The protein exhibits cytokine, cell differentiation activities and induces production of cother cytokines in certain cell populations. The protein also exhibits cother cytokines in certain cell populations. The protein also exhibits cother cytokines in certain cell populations. The protein also exhibits cother cytokines in certain cell populations. The protein also exhibits cother cytokines in certain cell populations. The protein also exhibits cother cytokines in certain cell populations. The protein also exhibits cother cytokines such as asthman deficiencies and disorders, and to treat cell conditions such as asthman, and in immune suppression after organization. The protein is useful in regulation of haematopolesis can deficiencies. It is also used in compositions for tissue growth or cellicatencies. It is also used in compositions for tissue growth or consequently in the treatment of periodontal disease and other coth repair processes. The protein is used in the treatment of osteoporosis cor osteoarthritis and in the treatment of periodontal disease and other coth repair processes. They are useful for protection or regeneration and contraceptives since they exhibit activin or inhibin related activities and as fertility inducing therapeutic. They are used for treatment of or processes. They are useful ferentiation. They are used for canditions resulting from coagulation activities e.g. myocardial confictions or agonists of receptor/ligand interactions. They are used to capacity themselves of receptor ligands or capacity, arthritis, and nephritis. They can be used to This sequence encodes a human protein of the invention, which has

Sequence 467 BP; 118 P 101 Ç 121 G G Η.

Alignment Scores

Qy 81 Db 354	Qy 61 Db 294	Qy 41 Db 234	Qy 21 Db 174	Qy . 1 Db 114	US-09-801-115B-2	Pred. No.: Score: Percent Similarity: Best Local Similarity: Query Match: DB:
PheAsnProSer	AlaLeuValThr	ValSerValLeu GTATCTGTGTTG	VallysMetLeu GTGAAGATGCTG	MetAspAsnVal arggaraacgro	(1-99) x	
GlyProTyrGlnL GGTCCTTACCAGC	AlaValCysCysL CAGTATGCTGTC	AlaLeuIleProG CACTGATACCAG	ArgLeuAspIleI cgcTgGATATTA	GlnProLysIleL CAGCCGAAAATAA	AAA15929 (1-467)	2.19e-57 504.00 100.00% 98.99% 99.21% 21
ysLysProValHi :: aaaagccTgTgCa	euAlaAspGlyAla TTGCCGACGGGGC	luThrThrThrLev aaaccacaacatto	leAsnSerLeuVa CAACTCACTGGT	ysHisArgProPho hACATCGCCCCTTO	67)	Length: Matches: Conservative: Mismatches: Indels: Gaps:
PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99 	AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu	ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyValPhe	VallysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle	MetAspasnvalGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis 		467 98 0
0	353	60 293	40 233	20 . 173		

Search completed: January 31, 2004, 04:20:24 Job time : 249 secs

GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.

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Run
                                                                                              OM protein - nucleic search, using frame_plus_p2n model
                                                 on:
January 31, 2004, 04:09:48; Search time 2058 Seconds (without alignments) 1169.166 Million cell updates/sec
                                                                                                                                                                Copyright
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Title: Perfect score: Scoring table: Sequence: Xgapop 10.0 , Xgapext Ygapop 10.0 , Ygapext Fgapop 6.0 , Fgapext Delop 6.0 , Delext US-09-801-115B-2 508 BLOSUM62 MDNVQPKIKHRPFCFSVKGH.....LFNPSGPYQKKPVHEKKEVL 0.5 7.0 7.0

Searched: 22781392 seqs, 12152238056 residues

Minimum DB seq length: 0
Maximum DB seq length: 200000000 Total number of hits satisfying chosen parameters:

45562784

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Command line parameters:

-MODELL-frame+ p2n.model -Dev=xlp
-MODELL-frame+ p2n.model -Dev=xlp
-Q2-(cgn2 1/USPTO_spool_p/US09801115/runat_30012004_113848_7035/app_query.fasta_1.263
-Q2-(cgn2 1/USPTO_spool_p/US09801115/runat_30012004_113848_7035/app_query.fasta_1.263
-DB=EST -QFWT=fastap -SUFFIX=p2n.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFWT=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=US09801115_0CGN_1 _3549_grunat_30012004_113848_7035 -NCPU=6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database : em_esthum: *
em_estin: * em_estba:*

222098765432 em_estro:
em_htc: *
gb_est1: *
gb_est2: *
gb_est3: * em_estov:*
em_estpl:* em_estmu:*

em_gss_mam:*
em_gss_mus:*
em_gss_pro::*
em_gss_phg:*
em_gss_vrl:*
gb_gssl:* gb_est4:*
gb_est5:* em_gss_vrt:* em_gss_fun:* em_gss_pln:* em em_gss_hum:* em_estom: * em_estfun:* _gss_inv:*

gb_gss2:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

REFERENCE	ORGANISM	KEYWORDS	ACCESSION	DEFINITION	LOCUS	AA911088/c	RESULT 1
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 386)	Homo sapiens	EST.	AA911088 1 GT-3050378	<pre>s1 NCI_CGAP_GC4 sence.</pre>	AA911088 386 bp mRNA linear EST 09-JUN-1998		

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ACCESSION
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                                     DEFINITION
                                                                         AI989739/c
                                                                                             RESULT 2
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Best Local Similarity:
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TITLE
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             391 b
wu21e10.x1 Soares_Dieckgraefe_
IMAGE:2520714 3', mRNA sequenc
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone lib="NCI_CGAP_GC4"
/clone lib="NCI_CGAP_GC4"
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo (dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="IMAGE:1519032"
/tissue_type="pooled germ
/lab_host="DH10B"
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                   mRNA sequence.
                           391 bp mRNA linear EST 27-OCT-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Butheria; Primates; Catarrhini; Hon
1 (bases 1 to 391)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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EST.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                 GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGGTGTTT
                                                                         AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
                                                                                                                                  ValSerValLeuAlaLeuIleProGluThrThrThrThrLeuThrValGlyGlyValPhe 60
                PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                        GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGCCCCTTATTTACCGGAAGCTTCTG
113
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="colonic mucosa from 3
disease"
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/db_xref="taxon:9606"
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 396)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
                                                                                      AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
                                                                                                                                                                ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                            GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                                                                   ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                                                                                                                                                                                                          ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
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//Clone lib="Soares_NFL_T_GBC_S1"
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//Clone lib="Soares_NFL_T_GBC_S1"
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//Clone lib="Soares NFL_T_GBC_S1"
//Clone lib="Soares NFL_T_GBC_S1"
a modified polylinker; Site_1: Not I; Site_2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHL19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-302087, 682632-687239,
726408-728711, and 729996-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
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/clone="IMAGE:2662317"
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                    41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                  21 VallysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arraying: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Cente Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html Insert Length: 527 Std Error: 0.00 Seq primer: -40ml3 fwd, ET from Amersham.
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ne58a03.s1 NCI_CGAP_Co3 Homo
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Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
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National Cancer Institute, Cancer Genome Anat
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GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
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/note="Vector: pT7T3D-Pac (Pharmacia) with a modified /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; lst strand cDNP was prepared from 12 pooled bulk tumor samples and primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                       normalization.
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/db_xref="taxon:9606"
/clone="IMAGE:901516"
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21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
                                                                                                                                           1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis 20
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Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40UP from Gibco.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI989747

wu21f09.x1 Soares_Dieckgraefe_cc
IMAGE:2520713 3', mRNA sequence
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1 (bases 1 to 402)

NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
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90 c 82 g 117 t
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/db_xref="taxon:9606"
/clone="IMAGE:2520713"
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08-09-801. Qy	egraped.	BASE COUNT	EATURES	REFERENCE AUTHORS TITLE JOURNAL COMMENT	RESULT 6 BM553628 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM	OY OY	Q B Q B
301-1158-2 (1-99) x bm555628 (1-904) 1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis	nt Scores: o.: Similarit cal Simila atch:	/organism="lone sapiens" //mol_type="mRNA" //db xref="taxon:9606" /clone="IMAGE:5740478" /tissue_type="duodenal adenocarcinoma, cell line" /tissue_type="duodenal adenocarcinoma, cell line" /lab host="DH10B (phage resistant)" /clone_lib="NH1M_MGC_88" /note="Organ: small intestine; Vector: pCMV-SPORTO /note="Organ: small intestine; Vector: pCMV-SPORTO Site_l: Nutl; Site_2: Sall; Cloned unidirectionall oligo-dT primed. Average insert size 1.767 kb. Lit enriched for full-length clones and constructed by Technologies. Note: this is a NIH_MGC Library." 121 a 80 c 87 g 116 t	Email Tissu CDNA CDNA CDNA CDNA DNA Clon found http: Plate High	Eukaryota; Metazoa; Chordata; Craniata; Vertek Mammalia; Eutheria; Primates; Catarrhini; Homi CE 1 (bases 1 to 404) RS NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene AL Unpublished Contact: Robert Strausberg, Ph.D.	BM553628 404 bp mRNA AGENCOURT_6541772 NIH_MGC_88 Homo sapien 5', mRNA sequence. BM553628 BM553628 1 GI:18792524 EST. Homo sapiens (human) Homo sapiens	188 GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG 81 PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99	
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 462 Std Error: 0.00
Seq primer: -40Up from Gibco.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Michael J. Brownstein,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AI826623 409 bp mRNA linear EST 21-DEC-1999 wk35610.x1 NCI_CGAP_Pr22 Homo sapiens cDNA clone IMAGE:2417395 3',
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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CDNA Library Arrayed by: Greg Lemon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG
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with a modified polylinker; 1st strand cDNA was prepared
from normal prostate bulk tissue, and was then primed with
a Not I - oligo(dT) primer. Double-stranded cDNA was
ligated to Eco RI adaptors (Pharmacia), digested with Not
I and cloned into the Not I and Eco RI sites of the
modified pT7T3 vector. Library is normalized, and was
constructed by Bento Soares and M. Fatima Bonaldo. "

94 c 80 g 123 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                             'clone
                                                                                                                                                                                                                                                                                                                                                                           tissue_type="normal prostate"
| host="DH10B"
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3.14e-56
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AI128804.1 GI:3597318
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                               normalization to a Cot = 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal lung library, Soares fetal lung
                                                                                  /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                  clone="IMAGE:1694391"
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                                                                                                                                                                                                                                                                                                                                                                                                                 Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R.
WashU-Merck EST Project 1997
                                                                                                                                                                                                                       IMAGE Consortium (info@image.llnl.gov) for Seq primer: -41ml3 fwd. ET from Amersham High quality sequence stop: 395.

Location/Qualifiers
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aa04a07.sl Soares_NhHMPu_Sl
                                                                                                                                                                                                                                                                                    Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL;
                                                                                                                                                                                                                                                                                                                                                 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                       Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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llier,L., Allen,M.,
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314 286 1810
/clone_lib="Soares_NhHMPu_S1"
/note="Organ: mixed_(see below); Vector: pT7T3D-Pac
/note="Organ: mixed a modified polylinker; Site_1: Not I;
Site_2: Eco RI; Equal amounts of plasmid DNA_from three
normalized libraries (melanocyte 2NbHM, pregnant uterus
                                                                                /clone="IMAGE:812244"
/tissue_type="Pooled human melanocyte,
pregnant_uterus"
/lab_host="DH10B"
                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:6043155"
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1 (bases 1 to 432)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore, K., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wyli,T., Waterston,R. and Wilson,R.

WashU_Merck_EST Project 1997
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zw67f10.sl Soares_testis_NHT
3', mRNA sequence
                                                                            Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -41ml3 fwd. ET from Amersham
High quality sequence stop: 421.
                                                                                                                                                                             4444 Forest Park Parkway, Box 8501, St. Louis,
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                         Contact: Wilson RK
Washington University School of Medicine
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                                           1. .432
/organism="Homo sapiens"
/mol_type="mRNA"
                                                             location/Qualifiers
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RESULT 11
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                                                                                                          COnLett: Noverthern Contact the Email: cgapbs-remail.nih.gov

Email: cgapbs-remail.nih.gov

This clone is available royalty-free through LLNL; contact the This clone is available royalty-free through LLNL; contact the This clone is available royalty-free through LLNL; contact the This clone is available royalty-free through LLNL; contact the This clone is a value of the This clone is value of the This clone is a value of the This clone is a value 
                       IMAGE Consortium (info@image.llnl.gov) fo
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 410.
Location/Qualifiers
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1 (bases 1 to 452)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/clone="IMAGE:781291"
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RESULT 12
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                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 453)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                   453 bp
wg90a02.xl Soares NSF F8 9W OT I
IMAGE:2372330 3', mRNA seguence
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AI743235.1 GI:5111523
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/clone lib="Soars NhHMPu S1"
/note="Organ: mixed (see Below); Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: Not I;
Site_2: Eco RI, Equal amounts of plasmid DNA from three
normalized libraries (melanocyte 2NbHM, pregnant uterus
NbHPU, and fetal heart NbHH19W) were mixed, and ss circles
were made in vitro. Following HAP purification, this DNA
was used as tracer in a subtractive hybridization
reaction. The driver was PCR-amplified cDNAs from pools of
5,000 clones made from the same 3 libraries. The pools
consisted of I.M.A.G.E. clones 260232-265223,
340488-345479, and 484488-489479."
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pregnant uterus"
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                                                                                                                                        466 bp ragencourt 10371072 NIH MGC 141 Homo IMAGE:6602806 5', mRNA sequence.
              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 466)
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Seq primer: -40UP f
NIH-MGC http://mgc.nci.nih.gov/
                                                                 Homo sapiens
                                                                                                                     BU564679.1 GI:22914979
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                                                                                 Homo sapiens (human)
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//Clone lib="Soares NSF F8 9W OT PA P S1"
//Clone lib="Soares Politication: pp7773D-Pac (Pharmacia) with
//note="Organ: pooled; Vector: pp7773D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site 2: Eco RI;
Equal amounts of plasmid DNA from five normalized
libraries were mixed, and ss circles were made in vitro.
Following HAP purification, this DNA was used as tracer in
subtractive hybridization reaction. The driver was
PCR-amplified cDNAs from pools of 5,000 clones made from
the same 5 libraries. The pools consisted of the following
libraries and cloneIDs: Soares NbHSP pool 1:
309384-310919, 323208-325895 Soares NbHSP pool 1:
145032-147335, 147720-148103, 148872-149255, 15002 -
150407, 151176-155327 Soares NbHFB-9W pool 1:
758280-760583, 772104-774407 Soares NbHBA pool 1:
304776-306311, 320136-322823, 326280-32663 Soares NbHOT
pool 1: 733720-726407, 739080-740999 Subtraction by Bento
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db_xref="taxon:9606"
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81 PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                                                                                                                                                                    41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
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Tissue Procurement: NCI
                                                                                                                     AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                             GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC 129
                                                                                 GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG
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/tissue type="mixed (pool of 40 RNAs)"
/tissue type="mixed (pool of 40 RNAs)"
/clone lib="NIH MGC 141"
/clone lib="NIH MGC 141"
/note="Wector: pDNR-LLB; Site 1: Sfil (ggccattatggcc);
/note="Wector: pDNR-LLB; Site 1: Sfil (ggccattatgcc);
/site_2: Sfil (ggccgcctggcc); Double-stranded cDNA was prepared from a pool of 40 cell line polyA+ RNAs (bladder - 2*, blood - 33.4*, brain - 5.6*, breast - 12.5*, colon - 4*, connective tissue - 1.4*, eye - 1*, intestine - 2.6*, kidney - 2.2*, liver - 5.7*, lung - 10.8*, NK-cell - 5.2*, ovary - 4*, pharynx - 2.5*, prostate - 4.3*, salivary gland - 1.3*, and skin - 2.3*). 5' and 3' adaptors were used in cloning as follows:
-AACCAGTGCTATCAACGCCGAAGTGGCCCATTACGGCCGG-3' and 5'-AACCAGTGGTATCAACGCAAGTGGCCCATTACGGCCGG-3' and 5'-AACCAGTGGTATCAACGCAAGTGGCCCATTACGGCCGGG-3' and 5'-AACCAGTGGTATCAACGCAAGTGGCCCATTACGGCCGGG-3' and 5'-AATTCTAGAGCCGAGGGCCGACATG-dT(30)NN-3'. Full-length enriched library was constructed using the Clontech Creator SMART kit and size-selected to contain the 0.2-0.5 kb size fraction (other fractions present in NIH MGC 142). Library created in the laboratory of M. Brownstein (NIMH, NIH). Note: this is a NIH_MGC Library."
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NIH).
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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468 bp mRNA linear EST 20-OCT-200
7171f03.x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone
IMAGE:3526805 3' Similar to TR:Q9UI41 Q9UI41 CHEMOKINE-LIKE FACTOR
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National Cancer Institute, Cancer Genome Anatomy
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    GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG
                           AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                          GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
                                                                                                                 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                                                                                                   MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 468)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Scares NDSHP pool 1: 309384-310915, 322208-225895 Scares NDSHP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002-150407, 151176-152327 Scares NDSHPR-W pool 1: 758280-760583, 772104-774407 Scartes NDSHPR pool 1: 304776-306311, 320136-322833, 32280-32663 Scares NDHOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Soares and M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pool 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="Soares NSF F8 9W OT PA P S1"
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site 2: Eco RI;
Equal amounts of plasmid DNA from five normalized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone="IMAGE:3526805"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lab_host="DH10B"
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508.00
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nd M. Fatima Bonaldo."
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Matches:
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REFERENCE
AUTHORS
TITLE
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SOURCE
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BF399486/c
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                                                          Best Local Similarity:
Query Match:
  US-09-801-115B-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURES
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                                                                                                   Percent Similarity:
                                                                                                                                                             Alignment Scores:
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PUBMED
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                                                                                                                                           No.:
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J75 Newton Road, 4156 MEBRF, Iowa City, IA 52242, USA

Tel: 319 335 8250

Fax: 319 335 9565

Email: bento-soares@uiowa.edu

The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonatide poly A tail. cDNA Library Preparation: W.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UI-R-CA1-bjb-b-12-0-UI.s1 UI-R-CA1 Rattus
UI-R-CA1-bjb-b-12-0-UI 3', mRNA sequence.
BF399486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: M13 Forward POLYA=Yes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Coordinated Laboratory for Computational Genomics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Soares, MB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genome Res. 6 (9), 791-806 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              discovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Normalization and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bonaldo, M.F.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 510)
                                                                                                                                                                                                                           130
    (1-99)
                                                                                                                                                                                                                                                         /note=Tvector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I, Site 2: Eco RI; The UI-R-CAI library is a subtracted library derived from the following tissues: thalamus, cerebellum, hypothalamus, medulla, pons, midbrain, cerebral cortex, corpus striatum, testis, and hippocampus. For a detailed description of the library from which this colone was derived, please visit our web site at ratest eng.uiowa.edu. The subtraction has been previously described in (Bonaldo, Lennon and Soares, Genome Research 6:791-806, 1996)
                                                                                                                                                                                                                           TAG_SEQ=None found"
131 c 104 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="UI-R-CA1-bjb-b-12-0-UI"
/lab_host="DH10B (Life Technologies)"
/clone_lib="UI-R-CA1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Rattus norvegicus"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /strain="Sprague-Dawley"
/db_xref="taxon:10116"
x BF399486
                                           4.14e-56
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    (1-510)
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Indels:
                                                                                                   Matches:
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1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis 20

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Ş	21	ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
Ъ	311	11 GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC 252
ΣΥ	41	ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe 60
8	251	GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT 192
Ωy	61	61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
В	191	GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG 132
δ	81	PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
용	131	TTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAA
Search completed: Ja	omple	Search completed: January 31, 2004, 05:25:28

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